

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
21 November 2002 (21.11.2002)

PCT

(10) International Publication Number
WO 02/093164 A2

(51) International Patent Classification⁷: **G01N 33/48**

(74) Agents: **LEIDESCHER, Thomas** et al.; Zimmermann & Partner, Postfach 330 920, 80069 München (DE).

(21) International Application Number: **PCT/EP02/05420**

(22) International Filing Date: **16 May 2002 (16.05.2002)**

(25) Filing Language: **English**

(26) Publication Language: **English**

(30) Priority Data:

01111858.5	16 May 2001 (16.05.2001)	EP
60/293,528	29 May 2001 (29.05.2001)	US
01117113.9	13 July 2001 (13.07.2001)	EP
60/305,898	18 July 2001 (18.07.2001)	US

(71) Applicant (for all designated States except US): **AXXIMA PHARMACEUTICALS AG** [DE/DE]; Am Klopferspitz 19, 82152 Martinsried (DE).

(72) Inventors; and

(75) Inventors/Applicants (for US only): **STEIN-GERLACH, Matthias** [DE/DE]; Stockdorfer Strasse 38A, 81475 München (DE). **SALASSIDIS, Konstadinos** [GR/DE]; Echinger strasse 20, 85386 Eching (DE). **BACHER, Gerald** [DE/DE]; Kriegerstrasse 62, 82110 Germering (DE). **MÜLLER, Stefan** [DE/DE]; Thalkirchner Str. 184, 81371 München (DE).

(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW.

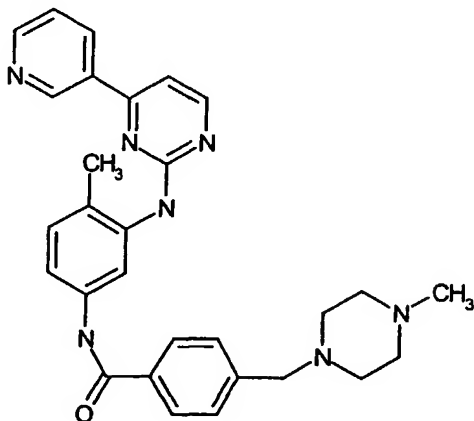
(84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Published:

— without international search report and to be republished upon receipt of that report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: **PYRIDYLPYRIMIDINE DERIVATIVES AS EFFECTIVE COMPOUNDS AGAINST PRION DISEASES**



Compound 53 (GleevecTM)

(57) Abstract: The present invention relates to pyridylpyrimidine derivatives of the general formula (I): wherein R represents hydrogen or methyl and Z represents nitrogen containing functional groups, the use of the pyridylpyrimidine derivatives as pharmaceutically active agents, especially for the prophylaxis and/or treatment of prion infections and prion diseases, as well as compositions containing at least one pyridylpyrimidine derivative and/or pharmaceutically acceptable salt thereof. Furthermore, the present invention is directed to methods for preventing and/or treating prion infections and prion diseases using said pyridylpyrimidine derivatives. Human cellular protein kinases, phosphatases and cellular signal transduction molecules are disclosed as targets for detecting, preventing and/or treating prion infections and diseases, especially BSE, vCJD, or CJD, which can be inhibited by the inventive pyridylpyrimidine derivatives.

WO 02/093164 A2

Pyridylpyrimidine derivatives as effective compounds against prion infections and prion diseases

5

Specification

The present invention relates to pyridylpyrimidine derivatives, the use of the pyridylpyrimidine derivatives as pharmaceutically active agents, especially for the prophylaxis and/or treatment of prion infections and prion diseases, as well as compositions containing at least one pyridylpyrimidine derivative and/or pharmaceutically acceptable salt thereof, and methods for preventing and/or treating prion infections and prion diseases. Furthermore, human cellular protein kinases, phosphatases and cellular signal transduction molecules are disclosed as targets for detecting, preventing and/or treating prion infections and diseases, especially BSE, vCJD, or CJD.

Background of the invention

Pyridylpyrimidine derivatives are known from WO 9509851 as effective compounds for chemotherapy of tumors, from WO 9509853, EP-A-0 588 762, WO 9509847, WO 9903854, and EP-B-0 564 409 as effective compounds for treatment of tumors. Furthermore, EP-B-0 564 409 discloses the use of said compounds in the treatment of arteriosclerosis and Exp. Opin. Ther. Patents, 1998, 8(12), 1599-1625 describes the use of pyridylpyrimidine derivatives, especially of GleevecTM, the Novartis compound CGP 57148, as tyrosine kinase inhibitors in cancer treatment.

Prions are infectious agents which do not have a nucleic acid genome. It seems that a protein alone is the infectious agent. A prion has been defined as "small proteinaceous infectious particle which resists inactivation by procedures that modify nucleic acids". The discovery that proteins alone can transmit an infectious disease has come as a considerable surprise to the scientific community. Prion diseases are often called "transmissible spongiform encephalopathies", because of the post mortem appearance of the brain with large vacuoles in the cortex and cerebellum. Probably most mammalian species develop these diseases. Prion diseases are a group of neurodegenerative disorders of humans and animals and the prion diseases can manifest as sporadic, genetic or infectious disorders. Examples for prion diseases acquired

by exogenous infection are the Bovine spongiform encephalitis (BSE) of cattle and the new variant of Creutzfeld-Jakob disease (vCJD) caused by BSE. Further examples include kuru, Gerstmann-Sträussler-Scheinker disease of humans as well as scrapie of animals. For many years, the prion diseases were thought to be caused by viruses despite intriguing evidence to the contrary. The unique characteristic common to all of these disorders, whether sporadic, dominantly inherited, or acquired by infection, is that they involve the aberrant metabolism of the prion protein (PrP). In many cases, the cellular prion protein (PrP^c) ["c" refers to cellular] is converted into the scrapie isoform (PrP^{Sc}) ["Sc" refers to Scrapie] by a posttranslational process that involves a conformational change. Often, the human prion diseases are transmissible to experimental animals and all of the inherited prion diseases segregate with PrP gene mutations.

These prion diseases in animals and humans have a long incubation period and a long clinical course, and are always fatal leading via decerebration to death within an average period of 7 months (CJD). Neuropathological features consist of neuronal vacuolization, neuronal death and gliosis with hyperastrocytosis. The precise diagnosis of transmissible neurodegenerative diseases can be established only by the examination of the central nervous system after biopsy or autopsy.

Clinical symptoms of the disease are progressive dementia, myoclonus and prominent ataxia with the additional clinical features of dysautonomia and delirious psychomotor excitement and with relatively preserved verbal responses.

Between 1980 and, roughly, 1996, about 750,000 cattle infected with BSE were slaughtered for human consumption in Great Britain (Anderson, R. M. *et al. Nature* 382, 779-788, 1996; Ferguson, N. M., Donnelly, C. A., Woolhouse, M. E. J. & Anderson, R. M. *Phil. Trans. R. Soc. Lond. B* 352, 803-838, 1997). The annual incidence of vCJD (3, 10, 10, 18, 14 and 33 deaths in 1995-2000, respectively) can be interpreted as a first sign of a steady or exponential increase over the next years. The suggestion by the European Union Scientific Steering Committee that up to 500,000 people could have been exposed to BSE from a single infected bovine has fuelled speculation that millions of consumers are at risk.

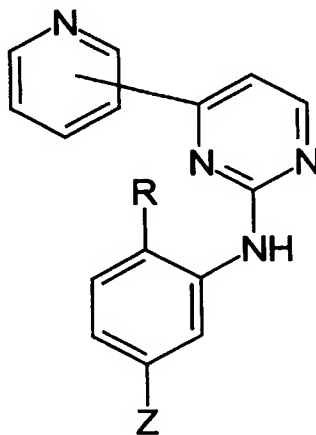
Recent findings demonstrate that the pathogenic PrP^{Sc} of vCJD can be found in the lymph system (e.g. tonsils, lymph nodes) in humans suggesting a high risk of horizontal spread via lymph and/or blood transmission, dramatically increasing the number of people at risk.

- The medical need in prion diseases today can be clearly defined as the establishment of a diagnostic system, that can detect the disease as early as possible in living humans and/or animals, to estimate the medical need for the treatment in the future and to identify the infected animals to remove them from the food chain. The medical need for prion diseases in the future (approximately starting in 5-10 years) will be medical treatment that inhibits the disease symptoms, the manifestation and/or progression of the disease.
- It is object of the present invention to provide novel and also known compounds which can be used as pharmaceutically active agents, especially for prophylaxis and/or treatment of prion infections and prion diseases, methods wherein said compounds are used in order to treat prion infections and prion diseases and compositions containing at least one inventive compound and/or pharmaceutically acceptable salt thereof as a pharmaceutically active ingredient.

The object of the present invention is solved by the teaching of the independent claims. Further advantageous features, aspects and details of the invention are evident from the dependent claims, the description, the examples, and the figures of the present application.

Description of the invention

One aspect of the present invention is related to compounds of the general formula (I):



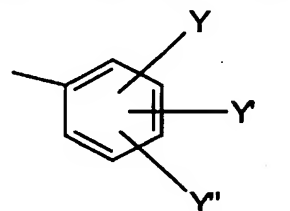
wherein:

R represents hydrogen or methyl;

Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

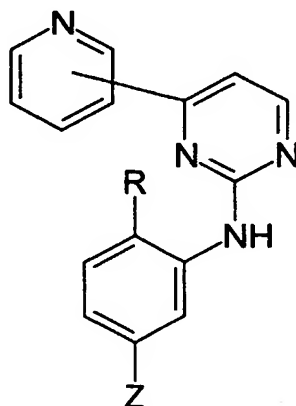
Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X,
5 -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl, cyclopentyl, pyridinyl, naphthyridinyl, or



and pharmaceutically acceptable salts thereof.

Another aspect of the present invention relates to the use of compounds of the
10 general formula (I):



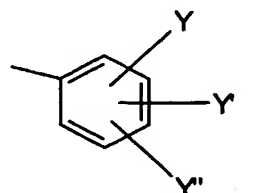
wherein:

R represents hydrogen or methyl;

Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-
15 methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl, cyclopentyl, pyridinyl, naphthyridinyl, or



and pharmaceutically acceptable salts thereof as pharmaceutically active agents, especially for prophylaxis and/or treatment of infectious diseases, or in a more general sense, for prophylaxis and/or treatment of neurodegenerative diseases.

- 5 Thus, one embodiment of the present invention disclosed herein is directed to a method for preventing and/or treating infections and/or diseases associated with said infections in an individual. Said method comprises administering to the individual an amount of at least one compound according to general formula (I) and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat
10 said infections and/or diseases. Most preferred is the administration of a compound 53.

As revealed for the first time herein, the present invention discloses the use of compounds of the general formula (I) for the prophylaxis and/or treatment of prion
15 infections and prion diseases. As described above, said pyridylpyrimidine derivatives have first of all been used in tumor therapy. The Novartis compound GleevecTM also known as GlivecTM, CGP-57148B, imatinib mesylate, STI-571, STI-571A, CAS 152459-95-5, or 4-((Methyl-1-piperazinyl)methyl)-N-[4-methyl-3-[[4-(3-pyridinyl)-2-pyrimidinyl]amino]-phenyl]benzamide methanesulfonate, has been
20 registered in many countries as anticancer drug. This GleevecTM compound (compound 53) is also the most active one in the indication prion diseases.

The name "prion" is used to describe the causative agents which underlie the transmissible spongiform encephalopathies. A prion is proposed to be a novel
25 infectious particle that differs from viruses and viroids. It is composed solely of one unique protein that resists most inactivation procedures such as heat, radiation, and proteases. The latter characteristic has led to the term protease-resistant isoform of the prion protein. The protease-resistant isoform has been proposed to slowly catalyze the conversion of the normal prion protein into the
30 abnormal form.

The term "isoform" in the context of prions means two proteins with exactly the same amino acid sequence that are folded into molecules with dramatically different tertiary structures. The normal cellular isoform of the prion protein
35 (PrP^C) has a high α -helix content, a low β -sheet content, and is sensitive to protease digestion. The abnormal, disease-causing isoform (PrP^{Sc}) has a lower α -helix content, a much higher β -sheet content, and is much more resistant to protease digestion.

Moreover, in a more general sense, the present invention is concerned with the prophylaxis and/or treatment of neurodegenerative diseases. For example, Alzheimer is a well-known neurodegenerative disease.

- 5 Preferred are the compounds wherein R represents hydrogen. Also preferred are compounds wherein Z represents $-\text{NH}-\text{CO}-\text{X}$ or $-\text{NH}-\text{SO}_2-\text{X}$ and/or wherein Y, Y', Y'' are independently of each other $-\text{H}$, $-\text{F}$, $-\text{Cl}$, $-\text{CH}_2\text{F}$, $-\text{CH}_2\text{Cl}$, $-\text{OH}$, $-\text{OCH}_3$, $-\text{CN}$, $-\text{OCF}_3$, or a 4-methylpiperazin-1-yl-methyl residue.
- 10 Also preferred are the following pyridylpyrimidine derivatives selected from the group comprising:
- Compound 1: (3-Nitrophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
 - Compound 2: (3-Aminophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
 - Compound 3: (5-Amino-2-methylphenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-
15 amine;
 - Compound 4: 4-Chloromethyl-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 5: 4-Chloromethyl-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - 20 Compound 6: 4-(4-Methylpiperazin-1-ylmethyl)-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 7: Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
 - Compound 8: 4-Chloro-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-
25 phenyl]-benzamide;
 - Compound 9: 4-Chloro-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 10: 3,4,5-Trimethoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - 30 Compound 11: 4-Cyano-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 12: 4-Methoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 13: 4-Chloro-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-
35 benzenesulfonamide;
 - Compound 14: Thiophene-3-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;

- Compound 15: 3,5-Dimethoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 16: 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 17: 4-Cyano-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 18: 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 19: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 10 Compound 20: Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 21: 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 Compound 22: 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 23: Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 24: Cyclohexanecarboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 20 Compound 25: Isoquinoline-5-sulfonic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 26: Isoquinoline-5-sulfonic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 Compound 27: (5-Nitro-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- Compound 28: (5-Amino-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- Compound 29: 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 30: 4-Cyano-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 31: (3-Aminophenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- Compound 32: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 35 Compound 33: Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 34: 4-Cyano-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;

- Compound 35: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 36: 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 37: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 38: Cyclohexanecarboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 39: 3,5-Dimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Compound 40: (5-Amino-2-methylphenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- Compound 41: Thiophene-3-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 15 Compound 42: 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 43: 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 44: (3-Aminophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- 20 Compound 45: (3-Nitrophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- Compound 46: 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 47: Isoquinoline-5-sulfonic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 Compound 48: 4-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 49: 4-Cyano-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 50: 3,4,5-Trimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 51: 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 52: 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 35 Compound 53: 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide (GleevecTM);
- Compound 54: 4-Methyl-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide

- Compound 55: 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 56: 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 57: Naphthalene-2-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 58: *N*-[3-(4-Pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 59: 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Compound 60: 4-Methoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 61: 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 62: Thiophene-2-carboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-15 amino)-phenyl]-amide;
- Compound 63: Naphthalene-2-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Compound 64: Isoquinoline-5-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- 20 Compound 65: Cyclopentanecarboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Compound 66: Naphthalene-2-carboxylic acid [3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 67: 4-Cyano-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-25 benzamide;
- Compound 68: 3,5-Dimethoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 69: 4-Bromo-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 70: 4-Methyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 71: 4-Fluoro-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 72: 3,5-Dichloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-35 benzamide;
- Compound 73: *N*-[3-(4-Pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 74: 4-Chloromethyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;

- Compound 75: 4-Methyl-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide
- Compound 76: 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 77: Naphthalene-2-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 78: 2-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 79: 2-Methoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Compound 80: 4-Methyl-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 81: 4-Methyl-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 Compound 82: *N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 83: 1-(3,5-Diacetyl-phenyl)-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- Compound 84: 1-{3,5-Bis-(amidinohydrazone)-phenyl}-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- 20 Compound 85: *N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- Compound 86: *N*-[3-(4-Pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- Compound 87: [1,8]Naphthyridine-2-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 Compound 88: [1,8]Naphthyridine-2-carbothioic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 89: 2-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 90: 4-Trifluoromethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 91: 4-Methyl-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;

35 and pharmaceutically active salts of these compounds.

Recent research has revealed how cells communicate with each other to coordinate the growth and maintenance of the multitude of tissues within the

human body. A key element of this communication network is the transmission of a signal from the exterior of a cell to its nucleus, which results in the activation or suppression of specific genes. This process is called signal transduction.

5 An integral part of signal transduction is the interaction of ligands, their receptors and intracellular signal transduction molecules. Ligands are messengers that bind to specific receptors on the surface of target cells. As a result of the binding, the receptors trigger the activation of a cascade of downstream signaling molecules, thereby transmitting the message from the exterior of the cell to its
10 nucleus. When the message reaches the nucleus, it initiates the modulation of specific genes, resulting in the production of RNA and finally proteins that carry out a specific biological function. Disturbed activity of signal transduction molecules may lead to the malfunctioning of cells and disease processes. Specifically, interference of the pathogenic PrP^{Sc} from prion diseases with
15 neuronal cells is necessary for the prion protein to induce its neuropathological features such as neuronal vacuolization, neuronal death and gliosis with hyperastrocytosis.

A key element of this communication network is the transmission of a signal from
20 the exterior of a cell to its nucleus, which results in the activation or suppression of specific genes. The human cellular protein kinases Abl and clk1 are two of the enzymes involved in said signal transduction process. As revealed herein said kinases Abl and clk1 serve as targets and are inhibited by the pyridylpyrimidine compounds of the general formula (I). It could be proved that prion infections
25 and/or prion diseases can be treated and also be prevented by the inhibition of said kinase Abl using the inventive pyridylpyrimidine derivatives. Inhibition of the kinase clk1 by said pyridylpyrimidine compounds can be used for the treatment of infections and diseases.

30 A microarray platform technology consisting of more than 1100 signal transduction cDNAs has been established. The technology is used for the identification of changes in RNA expression patterns as a result of the manipulation of the host cell by PrP^{Sc}. In addition, differential display techniques were used in order to pinpoint these changes to those enzymes which could be potential targets for drug
35 intervention.

Employing this predefined set of signal transduction relevant cDNAs on the filters, the expression pattern of signal transduction mRNAs in neuronal mouse cells

transfected with the pathogenic form of the prion protein (PrP^{Sc}) were compared with the same cells transfected with the non-pathogenic wild-type form (PrP^C) as a control. Interference of the PrP^{Sc} with the cellular signaling events is reflected in different gene expression when compared to the control cellular situation (PrP^C).

5

Using this technology, the human cellular protein kinases FGF-R1 (also known as flg, Fl-1, Flt-2, or b-FGFR), Tkt (also known as CCK-2, DDR-2, or EDDR, EC Number 2.7.1.112), Abl (also known as c-abl), clk1, MKK7 (also known as SKK4, SAPKK4, SAPKK5, or JNKK2), LIMK-2, CaM-KI, JNK2 (also known as SAPK1a, SAPKalpha), CDC2 (also known as CDK1), PRK, the human cellular protein phosphatases PTP-SL (also known as MCP83), PTP-zeta, the cellular signal transduction molecules HSP86, and GPIR-1 were identified as potential anti-prion disease targets. Said cellular protein kinases, phosphatases and signal transduction molecules are found to be specifically up- or downregulated by PrP^{Sc} in relevant mouse neuronal cells.

10

15

Surprisingly, it was found that the following human cellular targets are significantly up- or downregulated in prion infected cells:

20

<u>target</u>	<u>regulation</u>
FGF-R1	3.6 fold stronger
Abl	5.6 fold stronger
MKK7	4.1 fold stronger
CDC2	2.0 fold weaker
25 Tkt	2.1 fold stronger
LIMK-2	2.1 fold stronger
CaM-KI	2.1 fold stronger
JNK2	2.0 fold weaker
PRK	2.0 fold weaker
30 PTPzeta	4.6 fold weaker
PTP-SL	5.0 fold weaker
HSP86	4.1 fold weaker
GPIR-1	2.3 fold weaker

25

30

35

Thus, one aspect of the present invention relates to a method for preventing and/or treating prion infections and/or diseases associated with said prion infections in an individual which comprises administering to the individual an amount of at least one compound of the general formula (I) and/or pharmaceutically acceptable salts

thereof effective to prevent and/or treat said prion infections and/or prion diseases. Most preferred is the administration of a compound according to claim 8.

5 It could be proven that inhibition of one target selected from FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 was effective to treat prion diseases. Therefore, another aspect of the invention relates to a method for preventing and/or treating prion infections and/or prion diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one compound according of the
10 general formula (I) and/or pharmaceutically acceptable salts thereof which inhibits at least partially the activity of one target selectef from FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

15 The nucleoside sequences of the genes coding for the human cellular protein kinase Abl and the protein kinase clk1 and their amino acid sequences are disclosed in form of a sequence listing shown below. The nucleoside and amino acid sequences for the kinase Abl (Accession Number: M14752) and for the kinase clk1 (Accession Numbers: XM002520, NM004071, L29222, L29219) were obtained from NCBI (National Library of Medicine: PubMed).

20

The compounds of general formula (I) were identified as inhibitors of at least one target selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1 by the use of a method for detecting compounds useful for the prophylaxis and/or treatment of
25 prion infections and/or diseases. Said method comprises

- 30 a) contacting a test compound with at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1; and
b) detecting the activity of said human cellular protein kinase, phosphatase or cellular signal transduction molecule.

35

The activity of a human cellular protein kinase, phosphatase or cellular signal transduction molecule was preferably measured by means of an enzymatic assay.

As used herein, the term "inhibitor" refers to any compound capable of downregulating, decreasing, suppressing or otherwise regulating the amount and/or activity of at least one human cellular protein kinase, phosphatase or cellular

signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Generally, said inhibitors, including suicide inhibitors, may be proteins, oligo- and polypeptides, nucleic acids, genes, small chemical molecules, or other chemical moieties.

The present disclosure teaches for the first time the up- or downregulation of the above-mentioned human cellular protein kinases, phosphatases, or cellular signal transduction molecules specifically involved in prion infections and/or diseases. Thus, the present invention is also directed to a method for detecting prion infections and/or diseases in an individual comprising:

- a) providing a sample from said individual; and
- b) adding to said sample a pharmaceutically effective amount of at least one pharmaceutically active agent; and
- c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

As used herein the term "sample" refers to any sample that can be taken from a living animal or human for diagnostic purposes, especially said sample comprises blood, milk, saliva, sputum, excrement, urine, spinal cord liquid, liquor, lachrymal gland liquid, biopsies and all other samples that can be taken from a living animal or human for diagnostic purposes.

The term "individual" preferably refers to mammals, especially humans or ruminants. Ruminants are, for instance, muledeer, elk, cow, cattle, sheep, goat, deer, or buffalo. Minks are an example for mammals which do not belong to the species of ruminants.

As used herein the term "ruminants" refers to an animal, for instance, cattle, sheep, goat, deer, elk, or buffalo that has four separate stomach chambers, and is therefore able to digest a wide range of organic and plant foods. The term "ruminants" refers also to exotic ruminants, like captive nyala, gemsbok, Arabian oryx, eland, kudu, scimitar-horned oryx, ankole, or bison which are also accessible to develop spongiform encephalopathy.

A similar aspect of the present invention is directed to a method for detecting prion infections and/or prion diseases in cells, cell cultures and/or cell lysates comprising:

- a) providing said cells, cell cultures and/or cell lysates; and
- b) adding to said cells, cell cultures and/or cell lysates a pharmaceutically effective amount of at least one pharmaceutically active agent; and
- c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

Furthermore, it has been shown that the inhibition of at least one target selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1 has an effect on the production of prions. Therefore, another aspect of the invention relates to a method for regulating the production of prions in an individual or in cells comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

The inventive compounds according to general formula (I) are examples for the above-mentioned pharmaceutically active agent. Preferably the targets FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2 are used with said methods.

Another type of pharmaceutically active agents useful within the methods disclosed herein are monoclonal or polyclonal antibodies which bind to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Thus, a further aspect of the present invention is related to said monoclonal or polyclonal antibodies which bind to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1,

MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

Another embodiment of the present invention utilizes the scientific findings that
5 some targets such as JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 are downregulated during prion infection and that upregulation of the effected target by means of an activator leads to an alternative way of treating prion infections and diseases associated with prion infection.

10 Thus, a method was developed for regulating the production of prions either in an individual or in cells. Said methods comprise the step of administering an individual or the cells a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent activates at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction
15 molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or wherein said agent at least partially activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1,
20 MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

Preferably the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 are used within the above-described methods.

25 Because of the fact that the organism may upregulate a given target such as FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2 in order to compete with the prion infection, it is also a reasonable approach to further support said upregulation by means of an activator. Therefore, the above-mentioned methods
30 apply either to targets which are downregulated but also to targets which are upregulated.

The novel and partially known pyridylpyrimidine compounds of the general formula (I) represent a new class of pharmaceuticals highly useful for the prophylaxis and
35 treatment of prion infections and prion diseases.

Thus, a further aspect of the present invention describes the use of a compound of the general formula (I) and/or pharmaceutically acceptable salts thereof for the

manufacture of a pharmaceutical formulation for prophylaxis and/or treatment of prion infections and/or diseases induced or caused by prion infection.

5 As used herein the Term "prion diseases" refers to transmissible spongiform encephalopathies. This group of neurologic diseases affects humans and many species of animals causing a "sponge-like" degeneration of brain tissue. Among other unique features, all of these diseases are associated with the accumulation of an abnormal form of the prion protein in nerve cells that eventually leads to the death of the host. While prion diseases can all be transmitted from one host to another, it remains contentious as to whether a virus-like infectious agent or the abnormal prion protein itself, the prion, causes the conversion of normal to abnormal protein.

15 Probably most mammalian species develop prion diseases. Specific examples for animals include:

- **Scrapie** sheep, goat
- **TME** (transmissible mink encephalopathy): mink
- **CWD** (chronic wasting disease): muledeer, deer, elk
- **BSE** (bovine spongiform encephalopathy): cows, cattles

20

Humans are also susceptible to several prion diseases. Examples are:

- **CJD** Creutzfeld-Jacob Disease
- **GSS** Gerstmann-Sträussler-Scheinker syndrome
- **FFI** Fatal familial Insomnia
- **Kuru**
- **Alpers Syndrome**

25

30 The human prion diseases include kuru, sporadic Creutzfeldt-Jakob disease (sCJD), familial CJD (fCJD), iatrogenic CJD (iCJD), Gerstmann-Sträussler-Scheinker (GSS) disease, fatal familial insomnia (FFI), and, more recently, new variant CJD (nvCJD or vCJD). In addition to these human diseases, prion-related

diseases, have been recognized in several animal hosts. Scrapie is a naturally occurring disease of sheep and goats that causes ataxia, behavioral changes, and a severe pruritus that leads to scraping behavior, from which the disease was named. Additional prion diseases in animals include transmissible mink encephalopathy (TME), chronic wasting disease (CWD) of deer and elk, feline spongiform encephalopathy (FSE), and bovine spongiform encephalopathy (BSE), among others.

The transmissible nature of prion disease was first demonstrated experimentally in 1936 when Cuillé and Chelle transmitted scrapie to a healthy goat by the intraocular administration of scrapie-infected spinal cord. Thirty years later, sCJD was transmitted to chimpanzees. The pathologic feature common to all these diseases is a prominent vacuolation of the gray matter of the brain that produces a "sponge-like" appearance on light microscopy. This histopathologic appearance, coupled with the transmissible nature of these diseases, led to their collective designation as "transmissible spongiform encephalopathies" or TSEs.

The etiologic agent of the TSEs was proposed to be a "slow virus" to explain its transmissible nature and the prolonged incubation period observed during experimental transmission studies. Early experiments suggested that protein may be a critical component of the infectious agent. These studies established the basis for a new form of a transmissible pathogen, one that is composed ostensibly of only protein and lacks any replicative elements such as nucleic acid.

The term "prion" was coined to indicate an *infectious* agent with *proteinlike* properties. The unusual properties of the pathogen were demonstrated in early experiments in which conditions that degrade nucleic acids, such as exposure to ionizing and ultraviolet radiation, did not reduce the infectivity of scrapie fractions. On the other hand, treatments that degrade protein, such as prolonged exposure to proteases, correlated with a reduction in infectivity. A protein with relative resistance to protease digestion was found to be consistently present in the brains of animals and humans with TSE. Surprisingly, this protein was found to be one that is normally encoded by a chromosomal gene of the host.

Thus, the question raised, how a normally expressed protein could also be a transmissible pathogen? It was hypothesized and later demonstrated that PrP exists in two major isoforms: the nonpathogenic or cellular form, designated PrP^C, and the pathogenic or scrapie-inducing form, designated PrP^{Sc}. Both PrP^C and

PrP^{Sc} have the same amino acid sequence, yet they differ in their biochemical properties: PrP^C is soluble in nondenaturing detergents and completely degraded by proteases, whereas PrP^{Sc} is insoluble in nondenaturing detergents and shows a relative resistance to proteases. Structural studies of PrP^C and PrP^{Sc} indicate a difference in the conformation of the two isoforms: PrP^C is predominantly helical, whereas PrP^{Sc} contains at least 40% pleated sheet structure. Conversion to this sheet structure appears to be the fundamental event in prion disease. The ultimate mechanism of how cells die coincident with the generation of prions is still unclear. Simple accumulation of pathogenic protein may not be sufficient to explain disease, however, it may constitute a critical step in cellular dysfunction.

It was shown that the pyridylpyrimidine compounds of the general formula (I) are highly effective for the prophylaxis and/or treatment of prion infections and/or prion diseases selected from the group comprising Scrapie, TME, CWD, BSE, CJD, vCJD, GSS, FFI, Kuru, and Alpers Syndrome. Preferably, the pyridylpyrimidine derivatives are used for preventing and/or treating BSE, vCJD, or CJD.

The above-mentioned prion infections and/or diseases associated with prion infections can be treated using the inventive pyridylpyrimidine derivatives by targeting at least one of the human cellular protein kinases, phosphatases or cellular signal transduction molecules selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1. Thereby, the compounds according to general formula (I) act as inhibitors for at least one of the above-mentioned targets and especially as inhibitors for at least one enzyme selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2.

According to these findings a further aspect of the present invention is directed to a method for preventing and/or treating prion infections and/or prion diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

Another aspect is related to a method for preventing and/or treating prion infections and/or prion diseases in cells or cell cultures comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

The inventive pyridylpyrimidine compounds of formula (I) are examples for the above-mentioned inhibitor. Said pyridylpyrimidine compounds and/or pharmaceutically acceptable salts thereof are administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M, preferably in the range of 0.01 – 10 μ M, more preferably in the range of 0.01 – 1 μ M, and most preferably in the range of 0.01 – 0.1 μ M.

Because of the fact that the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 are downregulated in cells infected with prions, an upregulation of said targets represents another strategy in order to treat prion infections and diseases like CJD (nvCJD or vCJD) associated with prion infections. Said upregulation can be performed by activators.

An agent that is able to upregulate, increase, activate, or stimulate the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, but especially of JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 is named "activator".

Thus, another embodiment of the present invention describes a method for preventing and/or treating prion infections and/or diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which activates at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal

transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or which activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Preferably, said method is directed to the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

- 10 As used herein, the term "agent" or "pharmaceutically active agent" refers to any chemical compound capable of down- or upregulating, de- or increasing, suppressing, activation, stimulating or otherwise regulating the amount and/or activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Generally, said agents may be proteins, oligo- and polypeptides, nucleic acids, genes, aptamers, small chemical molecules, or other chemical moieties. An agent may be either an inhibitor or an activator and especially an inhibitor for the enzymes FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2 and an activator for the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

One special kind of said pharmaceutically active agents are aptamers which function as regulators of the activity of a wide range of cellular molecules such as human cellular protein kinase and phosphatase. Aptamers are nucleic acid molecules selected in vitro to bind small molecules, peptides, or proteins with high affinity and specificity. Aptamers not only exhibit highly specific molecular recognition properties but are also able to modulate the function of their cognate targets in a highly specific manner by agonistic or antagonistic mechanisms. Most famous examples for aptamers are DNA aptamers or RNA aptamers.

Further examples for pharmaceutically active agents are the pyridylpyrimidine compounds of the present invention and/or pharmaceutically acceptable salts thereof. Said compounds are administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M, preferably in the range of 0.01 – 10 μ M, more preferably in the range of 0.01 – 1 μ M, and most preferably in the range of 0.01 – 0.1 μ M.

The compounds of general formula (I) can be administered in a daily dosage in the range of 25 mg to 1000 mg, preferably in a daily dosage of 400 mg to 600 mg, more preferably in a daily dosage of 500 mg, and most preferably in continuously increased daily dosages starting at a initial daily dosage of 400 mg and ending up in a daily dosage of 600 mg at the end of the treatment.

A question is how PrP^C does convert to PrP^{Sc}? Potential mechanisms that initiate conversion of PrP^C to PrP^{Sc} include a germ line mutation of the human prion protein gene (PRNP), a somatic mutation within a particular neuron, and spontaneous conversion of PrP^C to an aberrant conformation that is not refolded appropriately to its native structure. The prion protein gene (PRNP) is the single gene on the short arm of chromosome 20 in humans which encodes the normal cellular isoform of the prion protein. Regardless of the initiating event, once an "infectious unit" has been generated, PrP^{Sc} appears to act as a conformational template by which PrP^C is converted to a new molecule of PrP^{Sc} through protein-protein interaction of PrP^{Sc} and PrP^C. This concept is supported by several studies which show that mice with the normal PrP gene deleted (PrP knockout mice) do not develop prion disease after inoculation with scrapie. Furthermore, transgenic (Tg) mice that express a chimeric PrP gene made of human and mouse segments develop protease-resistant chimeric mouse-human PrP^{Sc} in their brains when inoculated with brain extracts from humans with prion disease. These findings clearly illustrate that prions do not self-replicate but instead convert nonpathogenic PrP^C to pathogenic PrP^{Sc}.

In its sporadic or nonfamilial form, CJD is the most common of the human prion diseases. Confusion and forgetfulness which progress rapidly to severe cortical dementia in combination with ataxia, myoclonus, and an abnormal electroencephalogram (EEG) represents the "classic tetrad" of CJD. However, a host of other neurologic signs and symptoms, including diffuse or focal weakness, painful neuropathy, choreiform movements, hallucinations, cortical blindness, primary language disturbance, supranuclear ophthalmoplegia, and alien hand syndrome, among others, have been observed. As the disease progresses from the early stage, ataxia commonly limits the patient's mobility.

Familial CJD (fCJD) includes those cases with a dominantly inherited mutation of the PRNP gene, in which the pathologic features of spongiform change occur in the absence of GSS-type plaques. Although, familial cases of CJD tend to have a clinical and pathologic phenotype similar to that of sCJD.

The original description of a patient with the onset of ataxia and dysarthria followed by variable degrees of pyramidal and extrapyramidal symptoms and late developing dementia defines the classic presentation of **GSS**. The duration of said disease ranges from 2 to 10 years. Death usually results from secondary infection, often from aspiration pneumonia because of impaired swallowing. The presence of plaque deposits regionally or diffusely throughout the cortex that are immunoreactive to anti-human PrP antibodies is the hallmark of this form of prion disease.

FFI is a genetic disorder which manifests itself by many symptoms due to the degeneration of a certain part of the brain, the thalamus. The affected area of the brain is the area responsible for sleep, the thalamus. The thalamus is the center which communications from the brain to the body and the body to the brain pass through for proper directions to where a signal should be received. When sleep takes place, it is thought that the thalamus becomes less efficient at this signal transfer function allowing for the vegetative state of sleep to come over an individual. Consequently, the symptoms of fatal familial insomnia are directly related to the malfunction of the responsibilities of the thalamus, namely sleep.

There are four stages of the disease before an individual's life ends. The first stage is progressive insomnia, the characteristic feature of fatal familial insomnia. By now, there is no cure for this illness.

The term "familial" means: affecting several members of the same family, usually as a result of an underlying genetic mutation.

The occurrence of **vCJD** is sobering because it appears to represent a situation in which the prion has "jumped" species, in this case from cow to human. Because the pathologic features and clinical presentation of vCJD differ significantly from those of sCJD, it is considered a new "strain" of human prion disease. The same "protein signature" was observed following experimental transmission of BSE to several animal hosts, supporting the idea that vCJD results from the infection of humans with BSE. vCJD occurs primarily in younger individuals (average age 27) with a somewhat protracted course of approximately 16 months. The brain shows diffuse vacuolation and the presence of distinctive dense core PrP-containing plaques surrounded by a halo of spongiform change.

Kuru is the condition which first brought prion diseases to prominence in the 1950s. The disease was found in geographically isolated tribes in New Guinea. It was established that ingesting brain tissue of dead relatives for religious reasons was likely to be the route of transmission.

5

Alpers Syndrome is the name given to prion diseases in infants.

Scrapie is the accepted, albeit somewhat colloquial, name for the naturally occurring transmissible spongiform encephalopathy of sheep and goats found worldwide. Scrapie also infects laboratory mice and hamsters making it one of the most important sources of new scientific information about this group of disorders. Scrapie was the first example of this type of disease to be noticed and has been known about for many hundreds of years. There are two possible methods of transmission in sheep: a) Infection of pasture with placental tissue carrying the agent followed by ingestion, or b) direct sheep-lamb transmission.

15

CWD is a fatal neurodegenerative disease of deer and elk, now known to be a transmissible spongiform encephalopathy. To date, affected animals have been found exclusively in the United States.

20

BSE

Bovine spongiform encephalopathy or "mad cow disease" appears to have originated from scrapie that has been recognized in Europe since the mid-18th century. It has since spread to most sheep-breeding countries and is widespread in the United Kingdom, where until 1988 the rendered carcasses of livestock (including sheep) were fed to ruminants and other animals as a protein-rich nutritional supplement.

25

During rendering, carcasses from which all consumable parts had been removed were milled and then decomposed in large vats by boiling at atmospheric or higher pressures, producing an aqueous slurry of protein under a layer of fat (tallow). After the fat was removed, the slurry was desiccated into a meat and bone meal product that was packaged by the animal food industry and distributed to owners of livestock and other captive animals (e.g., zoo and laboratory animals, breeding species, pets).

30
35

A further aspect is related to a method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction

molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in an individual comprising the step of administering the individual a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent
5 inhibits at least partially the transcription of DNA or the translation of RNA.

And a still further aspect of the present invention relates to a method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1,
10 Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in the cells, the method comprising the step of administering the cells a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent inhibits at least partially the transcription of DNA or the translation of RNA.

15 As used herein, the term "regulating expression and/or activity" generally refers to any process that functions to control or modulate the quantity or activity (functionality) of a cellular component. Static regulation maintains expression and/or activity at some given level. Upregulation refers to a relative increase in
20 expression and/or activity. Accordingly downregulation refers to a relative decrease in expression and/or activity. Downregulation is synonymous with inhibition of a given cellular component's activity.

The transcription of DNA and the translation of RNA can be inhibited by
25 oligonucleotides or oligonucleotide derivatives. Thus, the present invention discloses oligonucleotides and derivatives of oligonucleotides which may be used in the above-mentioned methods. The oligonucleotide and/or its derivatives bind to the DNA and/or RNA encoding a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1,
30 Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 and suppress the transcription of DNA or translation of RNA.

As described above, said prion infection and/or disease associated with said prion infection is selected from the group comprising Scrapie, TME, CWD, BSE, vCJD, CJD, GSS, FFI, Kuru, and Alpers Syndrome. Preferably, the method is used for
35 prophylaxis and/or treatment of BSE, vCJD, or CJD. The above disclosed methods are preferably applied to CJD, vCJD, and BSE, more preferably applied to vCJD and BSE, and most preferably applied to BSE.

Some methods of the present invention identify compounds useful for prophylaxis and/or treatment of prion infections and/or diseases by screening a test compound, or a library of test compounds, for its ability to inhibit at least one of the above-mentioned human cellular protein kinases, phosphatases, or cellular signal transduction molecules, identified herein as characteristically up- or downregulated during prion production or growth inside a cell or individual. A variety of assay protocols and detection techniques are well known in the art and easily adapted for this purpose by a skilled practitioner. Such methods include, but are not limited to, high throughput assays (e.g., microarray technology, phage display technology), and *in vitro* and *in vivo* cellular and tissue assays.

Thus, a solid support is disclosed in the present invention useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized oligonucleotide, wherein said oligonucleotide encodes one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

A further aspect of the present invention is related to a solid support useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

In another embodiment, a component of the above-mentioned methods comprises peptide fragments of one or more of the above-identified human cellular protein kinases, phosphatases or cellular signal transduction molecules immobilized on a solid support. Once again the most preferred solid support embodiment would contain polymers of sufficient quality and quantity to detect all of the above-mentioned human cellular protein kinases, phosphatase and cellular signal transduction molecules (e.g., a nucleic acid or a peptide microarray). A variety of supports and constructions of the same for the methods disclosed herein are well known in the art and easily adapted for this purpose by a skilled practitioner (cf.,

for example: Marschall, 1999 "Do-it-yourself gene watching" Science 286, 444-447; Service 2000 "Protein arrays step out of DNA's shadow" Science 289, 1673).

It is preferred that mRNA is measured as an indication of expression. Methods for assaying for mRNA include, but are not limited to, Northern blots, slot blots, dot blots, and hybridization to an ordered array of oligonucleotides. Nucleic acid probes useful for assay of a sample are preferably of sufficient length to specifically hybridize only to appropriate, complementary transcripts. Typically the oligonucleotide probes will be at least 10 to 25 nucleotides in length. In some cases longer probes of at least 30 to 50 nucleotides will be desirable.

The cDNA oligonucleotides immobilized on said membrane filter which are used for detecting the up- or downregulation of the above-mentioned human cellular protein kinases, phosphatases, and cellular signal transduction molecules by hybridization to the radioactively labeled cDNA probes have the nucleotide sequences listed in table 1.

Table 1: Nucleotide sequences of cDNA-arrays

Cellular kinase, phosphatase, or signal transduction molecule	Sequence of immobilized DNA on arrays (in relation to the respective Acc No)
FGF-R1	41 bp – 2619bp (X52833)
Tkt (EC 2.7.1.112)	1 bp – 3096bp (X74764)
Abl	2153 bp – 3765 bp (M14752)
clk1	156 bp -1610 bp (L29219)
MKK7	77 bp – 1323 bp (AF013588)
CDC2	77 bp – 1050 bp (X05360)
CaMKI	145 bp – 1452 bp (L41816)
JNK2	507 bp – 1782 bp (L31951)
LIMK-2	963 bp – 2047 bp (D45906)
PRK	n.a bp – 1862 bp (U56998)
PTP zeta (EC 3.1.3.48)	148 bp – 7604 bp (X54135)
PTP-SL	862 bp – 1902 bp (NM_002849)
HSP86	n.a bp – n.a bp (X07270)
GPIR-1	n.a bp – n.a bp (n.a)

Tkt has been assigned to the EC Number: 2.7.1.112

PTP zeta has been assigned to the EC Number : 3.1.3.48

The nucleoside sequences of the genes coding for the human cellular protein kinases, phosphatases, or cellular signal transduction molecules listed in Table 1 together with the amino acid sequences and the enzyme commission numbers (E.C. numbers) of said enzymes can be obtained from NCBI (National Library of Medicine: PubMed; Web address: www.ncbi.nlm.nih.gov/entrez).

The polypeptide product of gene expression may be assayed to determine the amount of expression as well. Methods for assaying for a protein include, but are not limited to, western blot, immuno-precipitation, radioimmuno assay, and peptide immobilization in an ordered array. It is understood, however, that any method for specifically and quantitatively measuring a specific protein or mRNA product can be used.

A variety of supports upon which nucleic acids or peptides can be immobilized are known in the art, for example filters, or polyvinyl chloride dishes. Any solid surface to which oligonucleotides or peptides can be bound, either directly or indirectly, either covalently or non-covalently, can be used. A preferred solid support is a microarray membrane filter or a "biochip". These contain particular polymer probes in predetermined locations on the array. Each predetermined location may contain more than one molecule of the probe, but each molecule within the predetermined location has an identical sequence.

The present invention incorporates by reference in their entirety techniques well known in the field of molecular biology. These techniques include, but are not limited to, techniques described in the following publications:

Ausubel, F.M. et al. eds., "Short Protocols In Molecular Biology" 4th Ed. 1999, John Wiley & Sons, NY (ISBN 0-471-32938-X);

Old, R.W. & S.B. Primrose "Principles of Gene Manipulation: An Introduction To Genetic Engineering" 3rd Ed. 1985, Blackwell Scientific Publications, Boston. Studies in Microbiology: V.2, 409 pp. (ISBN 0-632-01318-4);

Mayer, R.J. & J.H. Walker eds. "Immunochemical Methods In Cell and Molecular Biology" 1987, Academic Press, London. 325 pp. (ISBN 0-12480-855-7);

Winnacker, E.L. "From Genes To Clones: Introduction To Gene Technology" 1987 VCH Publishers, NY. (translated by Horst Ibelgauf) 634 pp. (ISBN 0-89573-614-4).

As described above, a microarray platform technology was developed consisting of more than 1100 signal transduction cDNAs immobilized on a solid support. Thus, another aspect of the present invention is directed to a solid support useful for detecting prion infections and/or diseases in an individual, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

The present invention discloses also for the first time a solid support useful for detecting prion infections and/or diseases in cells, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

The present invention further incorporates by reference in their entirety techniques well known in the field of microarray construction and analysis. These techniques include, but are not limited to, techniques described in the following patents and patent applications describing array of biopolymeric compounds and methods for their fabrication:

U.S. Pat. Nos. 5,807,522; 6,087,102; WO 93/17126; WO 95/11995; WO 95/35505; EP 742 287; and EP 799 897.

Techniques also include, but are not limited to, techniques described in the following patents and patent application describing methods of using arrays in various applications:

U.S. Pat. Nos. 5,994,076; 6,033,860; 6,040,138; 6,040,140; WO 95/21265; WO 96/31622; WO 97/10365; WO 97/27317; EP 373 203; and EP 785 280

Still a further aspect of the present invention is directed to pharmaceutical compositions comprising at least one pharmaceutically active agent together with a pharmaceutically acceptable carrier, excipient or diluents. Examples for

pharmaceutically active agents are the above-mentioned inventive compounds according to formula (I), or other small chemical molecules, antibodies, aptamers, oligo- and polynucleotides, genes and other biological components capable of regulating the activity of at least one target selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or which are effective to treat prion infections and diseases associated with prion infection. Said prion infections and diseases are preferably Scrapie, TME, CWD, BSE, vCJD, CJD, GSS, FFI, Kuru, and Alpers Syndrome.

Thus, the pharmaceutical compositions according to the present invention may comprise an inhibitor, such as the inventive pyridylpyrimidine compounds or an activator such as aptamers for at least one target selected from FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. It is also possible to have a combination of inhibitors or activators as active ingredients in one single pharmaceutical composition. Furthermore, suitable are also combinations of at least one inhibitor and at least one activator for different targets within a single pharmaceutical composition. For example, a pharmaceutical composition could comprise compound 12 as an inhibitor for, for instance, the target Abl, and an activator such as an aptamer for, for instance, the human cellular protein kinase JNK2.

Said pharmaceutical compositions are useful for the prophylaxis and/or treatment of an individual afflicted with prions comprising at least one agent capable of inhibiting and/or activating at least partially the activity, the expression, and/or the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

The pyridylpyrimidine compounds of the present invention are basic and form pharmaceutically acceptable salts with organic and inorganic acids. Examples of suitable acids for such acid addition salt formation are hydrochloric acid, hydrobromic acid, sulfuric acid, phosphoric acid, acetic acid, citric acid, oxalic acid, malonic acid, salicylic acid, p-aminosalicylic acid, malic acid, fumaric acid, succinic acid, ascorbic acid, maleic acid, sulfonic acid, phosphonic acid, perchloric acid, nitric acid, formic acid, propionic acid, gluconic acid, lactic acid, tartaric acid, hydroxymaleic acid, pyruvic acid, phenylacetic acid, benzoic acid, p-aminobenzoic

acid, p-hydroxybenzoic acid, methanesulfonic acid, ethanesulfonic acid, nitrous acid, hydroxyethanesulfonic acid, ethylenesulfonic acid, p-toluenesulfonic acid, naphthylsulfonic acid, sulfanilic acid, camphorsulfonic acid, china acid, mandelic acid, o-methylmandelic acid, hydrogen-benzenesulfonic acid, picric acid, adipic acid, d-o-tolytartaric acid, tartronic acid, α -toluic acid, (o, m, p)-toluic acid, naphthylamine sulfonic acid, and other mineral or carboxylic acids well known to those skilled in the art. The salts are prepared by contacting the free base form with a sufficient amount of the desired acid to produce a salt in the conventional manner.

It is also possible to obtain acid addition salts with amino acids like methionine, tryptophane, lysine or arginine, especially with pyridylpyrimidine compounds of the general formula (I) carrying a carboxylic acid residue.

Depending upon the substituents on the inventive pyridylpyrimidine compounds, one may be able to form salts with bases, too. Thus, for example, if there are carboxylic acid substituents in the molecule, salts may be formed with inorganic as well as organic bases such as, for example, NaOH, KOH, NH_4OH , tetraalkylammonium hydroxide, and the like.

The compounds of the general formula (I) can also be administered in form of their pharmaceutically active salts optionally using substantially nontoxic pharmaceutically acceptable carriers, excipients or diluents. The medications of the present invention are prepared in a conventional solid or liquid carrier or diluents and a conventional pharmaceutically-made adjuvant at suitable dosage level in a known way. The preferred preparations are in administratable form which is suitable for oral application. These administratable forms, for example, include pills, tablets, film tablets, coated tablets, capsules, powders and deposits.

The preferred administratable forms are tablets, film tablets, coated tablets, gelatin capsules, and opaque capsules. Each pharmaceutical composition contains at least one compound of the general formula (I), preferably compound 53 and/or pharmaceutically acceptable salts thereof in an amount of 50 mg to 150 mg, preferably 80 mg to 120 mg, and most preferably in an amount of 100 mg per formulation.

Furthermore, the subject of the present invention also includes pharmaceutical preparations for parenteral, including dermal, intradermal, intragastrical,

intracutaneous, intravasal, intravenous, intramuscular, intraperitoneal, intranasal, intravaginal, intrabuccal, percutaneous, rectal, subcutaneous, sublingual, topical or transdermal application, which in addition to typical vehicles and diluents contain a pyridylpyrimidine compound of the general formula (I) and/or a pharmaceutically acceptable salt thereof as active ingredient.

Within the disclosed methods the pharmaceutical compositions of the present invention, containing pyridylpyrimidine derivatives of the general formula (I) as active ingredients, will typically be administered in admixture with suitable carrier materials selected with respect to the intended form of administration, i.e. oral tablets, capsules (either solid-filled, semi-solid filled or liquid filled), powders for constitution, oral gels, elixirs, dispersible granules, syrups, suspensions, and the like, and consistent with conventional pharmaceutical practices. For example, for oral administration in the form of tablets or capsules, the active drug component may be combined with any oral nontoxic pharmaceutically acceptable inert carrier, such as lactose, starch, sucrose, cellulose, magnesium stearate, dicalcium phosphate, calcium sulfate, talc, mannitol, ethyl alcohol (liquid forms) and the like. Moreover, when desired or needed, suitable binders, lubricants, disintegrating agents and coloring agents may also be incorporated in the mixture. Powders and tablets may be comprised of from about 5 to about 95 percent inventive composition.

Suitable binders include starch, gelatin, natural sugars, corn sweeteners, natural and synthetic gums such as acacia, sodium alginate, carboxymethylcellulose, polyethylene glycol and waxes. Among the lubricants, there may be mentioned for use in these dosage forms, boric acid, sodium benzoate, sodium acetate, sodium chloride, and the like. Disintegrants include starch, methylcellulose, guar gum and the like. Sweetening and flavoring agents and preservatives may also be included where appropriate. Some of the terms noted above, namely disintegrants, diluents, lubricants, binders and the like, are discussed in more detail below.

Additionally, the compositions of the present invention may be formulated in sustained release form to provide the rate controlled release of any one or more of the components or active ingredients to optimize the therapeutic effects, i.e. antihistaminic activity and the like. Suitable dosage forms for sustained release include layered tablets containing layers of varying disintegration rates or controlled release polymeric matrices impregnated with the active components

and shaped in tablet form or capsules containing such impregnated or encapsulated porous polymeric matrices.

5 Liquid form preparations include solutions, suspensions and emulsions. As an example may be mentioned water or water-propylene glycol solutions for parenteral injections or addition of sweeteners and opacifiers for oral solutions, suspensions and emulsions. Liquid form preparations may also include solutions for intranasal administration.

10 Aerosol preparations suitable for inhalation may include solutions and solids in powder form, which may be in combination with a pharmaceutically acceptable carrier such as inert compressed gas, e.g. nitrogen.

15 For preparing suppositories, a low melting wax such as a mixture of fatty acid glycerides such as cocoa butter is first melted, and the active ingredient is dispersed homogeneously therein by stirring or similar mixing. The molten homogeneous mixture is then poured into convenient sized molds, allowed to cool and thereby solidifies.

20 Also included are solid form preparations which are intended to be converted, shortly before use, to liquid form preparations for either oral or parenteral administration. Such liquid forms include solutions, suspensions and emulsions.

25 The inventive pyridylpyrimidine compounds of the present invention may also be deliverable transdermally. The transdermal compositions may take the form of creams, lotions, aerosols and/or emulsions and can be included in a transdermal patch of the matrix or reservoir type as are conventional in the art for this purpose.

30 The term capsule refers to a special container or enclosure made of methyl cellulose, polyvinyl alcohols, or denatured gelatins or starch for holding or containing compositions comprising the active ingredients. Hard shell capsules are typically made of blends of relatively high gel strength bone and pork skin gelatins. The capsule itself may contain small amounts of dyes, opaquing agents, plasticizers and preservatives.

35 Tablet means compressed or molded solid dosage form containing the active ingredients with suitable diluents. The tablet can be prepared by compression of

mixtures or granulations obtained by wet granulation, dry granulation or by compaction well known to a person skilled in the art.

5 Oral gels refers to the active ingredients dispersed or solubilized in a hydrophilic semi-solid matrix.

Powders for constitution refers to powder blends containing the active ingredients and suitable diluents which can be suspended in water or juices.

10 Suitable diluents are substances that usually make up the major portion of the composition or dosage form. Suitable diluents include sugars such as lactose, sucrose, mannitol and sorbitol, starches derived from wheat, corn rice and potato, and celluloses such as microcrystalline cellulose. The amount of diluents in the composition can range from about 5 to about 95% by weight of the total
15 composition, preferably from about 25 to about 75%, more preferably from about 30 to about 60% by weight.

The term disintegrants refers to materials added to the composition to help it break apart (disintegrate) and release the medicaments. Suitable disintegrants
20 include starches, "cold water soluble" modified starches such as sodium carboxymethyl starch, natural and synthetic gums such as locust bean, karaya, guar, tragacanth and agar, cellulose derivatives such as methylcellulose and sodium carboxymethylcellulose, microcrystalline celluloses and cross-linked microcrystalline celluloses such as sodium croscarmellose, alginates such as
25 alginic acid and sodium alginate, clays such as bentonites, and effervescent mixtures. The amount of disintegrant in the composition can range from about 2 to about 20% by weight of the composition, more preferably from about 5 to about 10% by weight.

30 Binders characterize substances that bind or "glue" powders together and make them cohesive by forming granules, thus serving as the "adhesive" in the formulation. Binders add cohesive strength already available in the diluents or bulking agent. Suitable binders include sugars such as sucrose, starches derived from wheat, corn rice and potato; natural gums such as acacia, gelatin and
35 tragacanth; derivatives of seaweed such as alginic acid, sodium alginate and ammonium calcium alginate; cellulosic materials such as methylcellulose and sodium carboxymethylcellulose and hydroxypropylmethylcellulose; polyvinylpyrrolidone; and inorganics such as magnesium aluminum silicate. The

amount of binder in the composition can range from about 2 to about 20% by weight of the composition, more preferably from about 3 to about 10% by weight, even more preferably from about 3 to about 6% by weight.

- 5 Lubricant refers to a substance added to the dosage form to enable the tablet, granules, etc. after it has been compressed, to release from the mold or die by reducing friction or wear. Suitable lubricants include metallic stearates such as magnesium stearate, calcium stearate or potassium stearate; stearic acid; high melting point waxes; and water soluble lubricants such as sodium chloride,
10 sodium benzoate, sodium acetate, sodium oleate, polyethylene glycols and D,L-leucine. Lubricants are usually added at the very last step before compression, since they must be present on the surfaces of the granules and in between them and the parts of the tablet press. The amount of lubricant in the composition can range from about 0.2 to about 5% by weight of the composition, preferably from
15 about 0.5 to about 2%, more preferably from about 0.3 to about 1.5% by weight.

- Glidants are materials that prevent caking and improve the flow characteristics of granulations, so that flow is smooth and uniform. Suitable glidants include silicon dioxide and talc. The amount of glident in the composition can range from about
20 0.1% to about 5% by weight of the total composition, preferably from about 0.5 to about 2% by weight.

- Coloring agents are excipients that provide coloration to the composition or the dosage form. Such excipients can include food grade dyes and food grade dyes
25 adsorbed onto a suitable adsorbent such as clay or aluminum oxide. The amount of the coloring agent can vary from about 0.1 to about 5% by weight of the composition, preferably from about 0.1 to about 1%.

- As used herein, a "pharmaceutically effective amount" of an inhibitor and/or an
30 activator is an amount effective to achieve the desired physiological result, either in cells treated *in vitro* or in a subject treated *in vivo*. Specifically, a pharmaceutically effective amount is an amount sufficient to inhibit and or activate, for some period of time, one or more of the clinically defined pathological processes associated with the prion infection. The effective amount may vary
35 depending on the specific inhibitor and/or activator selected, and is also dependent on a variety of factors and conditions related to the subject to be treated and the severity of the infection. For example, if an inhibitor and/or activator is to be administered *in vivo*, factors such as the age, weight and health

of the patient as well as dose response curves and toxicity data obtained in pre-clinical animal work would be among those considered. If the inhibitor and/or activator is to be contacted with the cells *in vitro*, one would also design a variety of pre-clinical *in vitro* studies to assess such parameters as uptake, half-life, dose, toxicity, etc. The determination of a pharmaceutically effective amount for a given pharmaceutically active agent is well within the ability of those skilled in the art.

It is also apparent to a person skilled in the art that detection includes any method known in the art useful to indicate the presence, absence, or amount of a detection target. Such methods may include, but are not limited to, any molecular or cellular techniques, used singularly or in combination, including, but not limited to: hybridization and/or binding techniques, including blotting techniques and immunoassays; labeling techniques (chemiluminescent, colorimetric, fluorescent, radioisotopic); spectroscopic techniques; separations technology, including precipitations, electrophoresis, chromatography, centrifugation, ultrafiltration, cell sorting; and enzymatic manipulations (e.g., digestion).

It should be stressed that all above-mentioned features, aspects, and details of the present invention discussed and described in connection with infections and infectious diseases, equally apply to neurodegenerative diseases, like Alzheimer.

It is readily apparent to those skilled in the art that other suitable modifications and adaptations of the compositions and methods of the invention described herein are evident and may be made without departing from the scope of the invention or the embodiments disclosed herein. Having now described the present invention in detail, the same will be more clearly understood by reference to the following examples, which are included for purposes of illustration only and are not intended to be limiting of the invention.

Description of figures

Fig. 1 shows 6 selected pyridylpyrimidine derivatives which are suitable inhibitors for prion diseases, namely compounds 4, 5, 37, 52, 84, and 88;

Fig. 2 shows the compound 4-(4-Methylpiperazin-1-ylmethyl)-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide, also known as GleevecTM;

Fig. 3 shows selected compounds that have been identified as potent inhibitors in a prion propagation assay at a concentration of 5 μ m.

Examples

Materials and methods

5

1. Generation of cDNA-arrays on membranes

In order to manufacture cDNAs-arrays on membranes, the following strategy was pursued: cDNAs encoding parts of or full length proteins of interest – in the following referred to as “target cDNAs” – were cloned into the plasmid Bluescript II KS⁺ (Stratagene, USA). Large scale purifications of these plasmids were performed according to standard techniques and 200 µl aliquots (1 µg/µl plasmid concentration) were transferred into appropriate 96well plates. Plates were closed with sealing tape and chilled on ice for 5 minutes after incubation for 10 minutes at 95°C. 10 µl of 0.6 N NaOH were added and the mix was stored for 20 minutes at room temperature before addition of 10 µl 2.5 M Tris-HCl pH 7.1 and 20 µl 40x SSC (3 M NaCl, 300 mM Sodium Citrate, pH 7.0). Target cDNAs were spotted onto Nylon or Nitrocellulose membranes using a BioGrid (BioRobotics, UK) equipped with a 0.7 mm pintool. In this way, between 200 ng and 350 ng of plasmids encoding target cDNAs were transferred onto the membranes and crosslinked to the membranes by ultraviolet light (1.2×10^5 µJ/cm²). The arrays were stored for use in subsequent experiments at room temperature.

25

2. Generation of cells

PrP^{Sc}- and PrP^C-transfected mouse neuronal cells (N2A) were cultured in MEM (Minimum Essential Medium, Life Technologies) supplemented with 10% fetal calf serum at 37°C and 5% CO₂ to obtain $\sim 6 \times 10^6$ cells per tissue culture flask.

30

3. Lysis of cells, isolation of total RNA and purification of polyA⁺ RNA

After incubation of the cells with the virus for the respective time-points, cells were washed twice with phosphate buffered saline (PBS) and then trypsinized. Subsequently, cells were removed from the culture dish by resuspension with PBS. Afterwards, cells were sedimented and directly lysed in Tri reagent by repetitive pipetting using in 1ml of Tri reagent (Molecular Research Centre, Inc., USA) per 1×10^6 cells.

The lysates were stored at room temperature for 5 minutes and then centrifuged at 12000xg for 15 minutes at 4°C. The supernatant was mixed with 0,1 ml of 1-bromo-3-chloropropane per 1 ml of Tri reagent and vigorously shaken. The suspension was stored for 5 minutes at room temperature and then centrifuged at
5 12000xg for 15 minutes at 4°C.

The colourless upper phase was transferred into new tubes, mixed with 5 µl of poly-acryl-carrier (Molecular Research Centre, Inc., USA) and with 0.5 ml of isopropanol per 1 ml of Tri reagent and vigorously shaken. The samples were
10 stored at room temperature for 5 minutes and then centrifuged at 12000xg for 8 minutes at 4°C. The supernatant was removed and the RNA pellet washed twice with 1 ml of 75% ethanol. The pellet was dried and resuspended for 10 minutes at 55°C in 50 µl of RNase-free buffer (5 mM Tris-HCl pH 7.5). The integrity of the isolated RNA was determined by agarose/formaldehyde gel electrophoresis and
15 the RNA was finally stored at -70°C for use in subsequent experiments.

4. Preparation of radioactively labelled cDNA probes from RNA

In order to obtain radioactively labelled cDNA probes total RNA was transcribed
20 into a cDNA-probe in the presence of radioactively labelled dATP. 12 µl bidistilled DEPC (Diethylpyrocarbonate) treated H₂O containing 0.5 µg of primer TXN (5'-TTT TTT TTT TTT TTT TXN-3' with T → dTTP; N → dATP, dCTP, dGTP or dTTP; X → dATP, dCTP or dGTP) and total RNA (1 to 10 µg) were shaken between 5 and 15' at 60°C and then incubated on ice for 2 minutes. After
25 centrifugation (30 seconds, 10000xg) 7 µl of a mix consisting of 100 µCi dATP-P³³ (Amersham, UK) which were dried under vacuum previously and resuspended in 4 µl first strand buffer (Life Technologies, USA), 2 µl 0.1M DTT (Dithiothreitol) and 1 µl labelling solution (4 mM dCTP, dGTP, dTTP each and 80 µM dATP final concentration) were added. Following the addition of 1 µl Superscript II reverse
30 transcriptase (Life Technologies, USA) the reaction was incubated for 10 minutes at room temperature and then for 60 minutes at 38°C. Subsequently, the reaction was vigorously shaken for 30 minutes at 68°C after adding 5 µl 0.5 M EDTA and 25 µl 0.6M NaOH.

35 Unincorporated nucleotides were removed from the labelling reaction using ProbeQuant G-50 columns (Amersham, UK). The column was vigorously shaken and centrifuged for 1 minute at 735xg in an appropriate reaction tube after bottom closure and lid were removed. The column was placed into a new reaction tube,

the probe was applied onto the centre of the column material and the column was centrifuged for 2 minutes at 735xg. The flow-trough was transferred into new reaction tubes and filled up to a volume of 100 µl with bidistilled H₂O. The probe was precipitated by centrifugation for 15 minutes at 12000xg after 4 µl 5M NaCl, 1 µl poly-acryl-carrier (Molecular Research Centre, Inc., USA) and 250 µl ethanol were added. The supernatant was discarded and the pellet was dried at 50°C for 5 minutes before starting with the hybridisation.

5. Hybridisation of radioactively labelled cDNA-probes to cDNA-arrays

The pellet was resuspended in 10 µl C₀T DNA (1 µg/µl, Roche Diagnostics, Germany), 10 µl yeast tRNA (1µg/µl Sigma, USA) and 10 µl polyA (1 µg/µl, Roche Diagnostics, Germany) and incubated at 55°C for 5 minutes. Herring sperm DNA was added to a final concentration of 100 µg/ml and the volume was filled up to 100 µl with 5 µl 10% SDS (Sodiumdodecylsulfat), 25 µl 20x SSPE (3M Sodium chloride, 0,2 M Sodium dihydrogen phosphate monohydrate, 0,02 M Ethylenedinitrilo tetraacetic acid, disodium salt dihydrate; pH 7,4) and bidistilled H₂O. The mix was put on 95°C for 5 minutes, centrifuged for 30 seconds at 10000xg and vigorously shaken for 60 minutes at 65°C. A 1 µl aliquot of the probe was used to measure the incorporation of radioactive dATP with a scintillation counter. Probes with at least a total of 20x10⁶ cpm were used.

The arrays were prehybridised for at least 3 hours at 42°C in hybridisation solution in a roller bottle oven. After prehybridization the radioactively labelled probe was added into the hybridisation solution and hybridisation was continued for 20 to 40 hours.

The probe was discarded and replaced with wash solution A (2xSSC). The arrays were washed twice in wash solution A at room temperature in the roller oven. Afterwards, wash solution A was replaced by wash solution B (2x SSC, 0.5% SDS) preheated to 65°C and arrays were washed twice for 30 minutes at 65°C. Then, wash solution B was replaced by wash solution C (0.5x SSC, 0.5% SDS) preheated to 65°C and arrays were washed twice for 30 minutes at 65°C. The moist arrays were wrapped in airtight bags and exposed for 8 to 72 hours on erased phosphoimager screens (Fujifilm, Japan).

6. Analysis of cDNA-arrays

The exposed phosphorimager screens were scanned with a resolution of 100 μ and 16bits per pixel using a BAS-1800 (Fujifilm, Japan). Files were imported into the computer program ArrayVision (Imaging Research, Canada). Using the program's features, the hybridization signals of each target cDNA were converted into numbers. The strength of the hybridization signals reflected the quantity of RNA molecules present in the probe. Differentially expressed genes were selected according to the ratio of their signal strength after normalization to the overall intensity of the arrays.

7. Cell culture and expression of 3F4-tagged PrP (3F4-ScN2a)

The mouse neuroblastoma cell line 3F4-ScN2a represents a stably transfected clone of ScN2a cells (PrP^{Sc} infected N2a cells) which overexpress 3F4-epitope-tagged murine PrP. Residues 109 and 112 of murine PrP were replaced by methionine to introduce the epitope for reactivity with the monoclonal anti-PrP antibody 3F4. Cells were maintained in Dulbecco's modified Eagle's (DMEM) or Opti-MEM medium containing 10 % fetal calf serum, antibiotics and glutamin. For generation of stable transfectants we used the vector pcDNA3.1/Zeo (Invitrogen; Leek, The Netherlands). Lipofection of cells with recombinant plasmids was done using standard procedures and recombinant clones were selected by addition of 300 μ g Zeocin/ml medium.

8. Treatment of cells with inhibitors

All tested compounds were solubilized in DMSO (dimethylsulfoxide), and prepared as 10 mM stock solutions. The drugs were applied to the cells described above for three days in final concentrations between 5 and 20 μ M.

9. Immunoblot and proteinase K (PK) analysis

Confluent cell cultures were lysed in cold lysis buffer (10 mM Tris-HCl, pH 7.5; 100 mM NaCl; 10 mM EDTA; 0.5 % Triton X-100; 0.5 % DOC) (EDTA: ethylene diamine tetraacetate; Triton X-100: t-octylphenoxypolyethoxyethanol; DOC: deoxycholic acid). Postnuclear lysates were split between those with and without proteinase K digestion. Samples without proteinase K digestion were supplemented with proteinase inhibitors (5 mM PMSF, 0.5 mM Pefabloc, and aprotinin) (PMSF: phenylmethylsulfonyl fluoride) and directly precipitated with ethanol. Samples for proteinase K digestion were incubated with 20 μ g/ml proteinase K for 30 min at 37°C; digestion was stopped with proteinase inhibitors,

and samples were ethanol precipitated. After centrifuging for 30 min at 3,500 rpm the pellets were redissolved in TNE buffer (10 mM Tris-HCl pH7.5, 100 mM NaCl, 1mM EDTA) and gel loading buffer was then added. After boiling for 5 min an aliquot was analyzed on 12.5 % PAGE. For Western blot analysis, the proteins
5 were electrotransferred to PVDF membranes (polyvinylidendifluoride). The membrane was blocked with 5 % non-fat dry milk in TBST (0.05 % Tween 20, 100 mM NaCl, 10 mM Tris-HCl, pH 7.8) (Tween 20: polyoxyethylenesorbitan monolaurate; Tris-HCl: Tris-(hydroxymethyl)-aminomethane-hydrochloride), incubated overnight with the primary antibody at 4°C and stained using the
10 enhanced chemiluminescence blotting kit from Amersham Corporation. Specific immuno-staining of the PrP^c and PrP^{Sc} forms were obtained with the prion protein specific antibody 3F4 (Signet Pathologies, U.S.A.).

10. Results

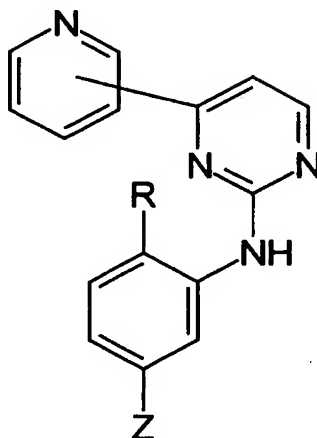
15 Determination of the amount of the pathogenic form of the prion protein PrP^{Sc} upon treatment of prion infected cells with different types of small molecule protein kinase inhibitors resulted in the identification of a compound class of pyridylpyrimidine derivatives exemplified by the compound 4-(4-Methylpiperazin-1-ylmethyl)-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide
20 (compound 53) and compounds 4, 5, and 37.

These compounds significantly reduced the amount of PrP^{Sc} in prion infected cells in a concentration range between 5 and 20 µM (final concentration). As shown in Fig. 3 the selected compounds 4, 5, 37, and 53 inhibit almost completely the
25 activity of prion propagation within said concentration range.

The compounds did not show any toxic effects on the cells in these concentrations. Therefore these molecules described herein serve as potential inhibitors for the medical intervention of prion diseases such as transmissible
30 spongiform encephalitis (TSE) infections which include Bovine spongiform encephalitis (BSE) or the new variant of Creutzfeld Jakob disease (vCJK).

Claims

1. Compounds having the general formula (I):



5

wherein:

R represents hydrogen or methyl;

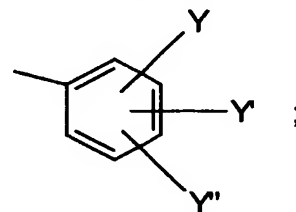
10

Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or

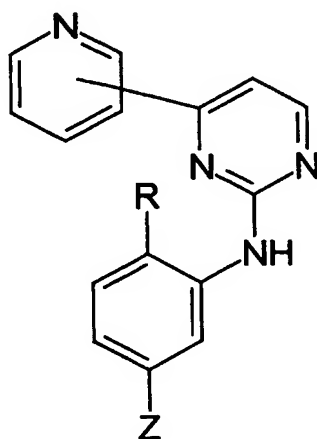


15

and pharmaceutically acceptable salts thereof.

20

2. Use of a compound having the general formula (I):



5 wherein:

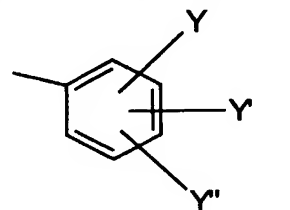
R represents hydrogen or methyl;

Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

10 Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or

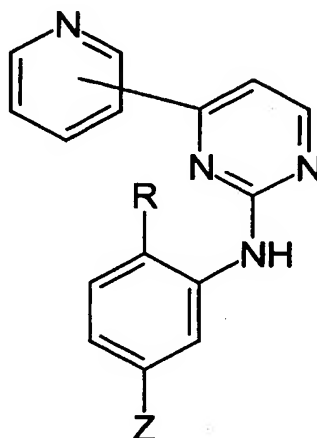


and pharmaceutically acceptable salts thereof as pharmaceutically active agents.

15

20

3. Use of a compound having the general formula (I):



5 wherein:

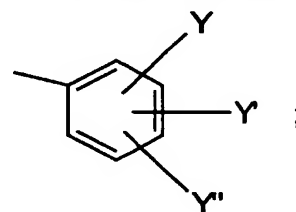
R represents hydrogen or methyl;

Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

10 Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or



and pharmaceutically acceptable salts thereof for prophylaxis and/or treatment of infectious diseases or neurodegenerative diseases.

15

4. Use of a compound according to claim 2 or 3 for the prophylaxis and/or treatment of prion infections and/or diseases induced by prion infection.

5. Use of a compound according to any one of claims 2 - 4 wherein R represents hydrogen.

20

6. Use of a compound according to any one of claims 2 - 5 wherein Z represents -NH-CO-X or -NH-SO₂-X.

7. Use of a compound according to any one of claims 2 – 6 wherein Y, Y', Y'' are independently of each other –H, –F, –Cl, –CH₂F, –CH₂Cl, –OH, –OCH₃, –CH₃, –CN, –OCF₃, 4-methylpiperazin-1-yl-methyl.
- 5 8. Use of a compound according to claim 2 or 3 wherein the compound is selected from the group comprising:
- (3-Nitrophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
(3-Aminophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
10 (5-Amino-2-methylphenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
4-Chloromethyl-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Chloromethyl-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-(4-Methylpiperazin-1-ylmethyl)-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-
15 phenyl]-benzamide;
Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
4-Chloro-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Chloro-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
20 3,4,5-Trimethoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Cyano-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Methoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
25 4-Chloro-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
Thiophene-3-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
3,5-Dimethoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
30 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Cyano-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-
35 benzamide;
4-Chloro-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;

- Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 10 Cyclohexanecarboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Isoquinoline-5-sulfonic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Isoquinoline-5-sulfonic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 15 (5-Nitro-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- (5-Amino-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Cyano-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 (3-Aminophenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- 4-Chloro-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 4-Cyano-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 30 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Cyclohexanecarboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 35 3,5-Dimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- (5-Amino-2-methylphenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;

- Thiophene-3-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 5 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- (3-Aminophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- (3-Nitrophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Isoquinoline-5-sulfonic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Cyano-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,4,5-Trimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-
- 15 benzamide;
- 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methyl-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-
- 25 benzamide;
- 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Naphthalene-2-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 30 *N*-[3-(4-Pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 35 Thiophene-2-carboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Naphthalene-2-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;

- Isoquinoline-5-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Cyclopentanecarboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- 5 Naphthalene-2-carboxylic acid [3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Cyano-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,5-Dimethoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Bromo-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 4-Methyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Fluoro-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 3,5-Dichloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- N*-[3-(4-Pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 4-Chloromethyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methyl-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 Naphthalene-2-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 2-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 2-Methoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 25 4-Methyl-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methyl-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 1-(3,5-Diacetyl-phenyl)-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- 30 1-{3,5-Bis-(amidinohydrazone)-phenyl}-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- N*-[3-(4-Pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- 35 [1,8]Naphthyridine-2-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- [1,8]Naphthyridine-2-carbothioic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;

2-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Trifluoromethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-
benzamide;
4-Methyl-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
5 and/or a pharmaceutically acceptable salt of these compounds.

9. Use according to claim 8 wherein the compound is 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide.

10. Use of a compound recited in any one of claims 2 – 9 and/or pharmaceutically acceptable salts thereof for the manufacture of a pharmaceutical composition for prophylaxis and/or treatment of prion infections and/or diseases induced by prion infection and/or neurodegenerative diseases.

11. Use according to claim 4 or 10 wherein said prion infection and/or disease is selected from the group comprising Scrapie, TME, CWD, BSE, CJD, vCJD, GSS, FFI, Kuru, and Alpers Syndrome.

12. Use according to claim 11 wherein said prion infection is BSE, vCJD, or CJD.

13. Use of a compound recited in any one of claims 2 – 9 as an inhibitor for at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

14. Use of a compound according to any one of claims 2 to 13 wherein the compound of the general formula (I) and/or pharmaceutically acceptable salts thereof is administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M.

15. Pharmaceutical composition comprising at least one compound recited in any one of claims 2 – 9 as an active ingredient, together with one or more pharmaceutically acceptable carrier(s), excipient(s) or diluents.

16. Method for preventing and/or treating infections and/or diseases in an individual which comprises administering to the individual an amount of at least one compound recited in claims 2 – 9 and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat said infections and/or diseases.
17. Method for preventing and/or treating prion infections and/or prion diseases induced by prion infections in an individual which comprises administering to the individual an amount of at least one compound recited in any one of claims 3 to 8 and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat said prion infection and/or disease.
18. Method for preventing and/or treating prion infections and/or prion diseases induced by prion infections in an individual which comprises administering to the individual an amount of at least one compound recited in claim 8 and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat said prion infection and/or disease.
19. Method for detecting prion infections and/or prion diseases in an individual comprising:
- a) providing a sample from said individual;
 - b) adding to said sample a pharmaceutically effective amount of at least one pharmaceutically active agent; and
 - c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.
20. Method according to claim 19 wherein said sample comprises blood, milk, saliva, sputum, excrement, urine, spinal cord liquid, liquor, lachrymal gland liquid, biopsies and all other samples that can be taken from a living animal or human for diagnostic purposes.
21. Method for detecting prion infections and/or prion diseases in cells, cell cultures and/or cell lysates comprising:
- a) providing said cells, cell cultures and/or cell lysates;
 - b) adding to said cells, cell cultures and/or cell lysates a pharmaceutically effective amount of at least one pharmaceutically active agent; and

- c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

5

22. Method for preventing and/or treating prion infections and/or prion diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

10

15

23. Method for preventing and/or treating prion infections and/or prion diseases in cell or cell cultures comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

20

25

24. Method for regulating the production of prions in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt,

30

35

Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

- 5 25. Method for regulating the production of prions in cells comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, 10 HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.
- 15 26. A monoclonal or polyclonal antibody that binds to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 20 27. Method according to any one of claims 19 – 25, wherein the agent is a monoclonal or polyclonal antibody which binds to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, 25 JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 30 28. Method according to any one of claims 19 – 25, wherein the agent is at least one compound of the general formula (I) and/or pharmaceutically acceptable salts thereof.
- 35 29. Method according to any one of claims 16 – 25, wherein the agent is 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-yl-amino)-phenyl]-benzamide and/or pharmaceutically acceptable salts thereof.
30. Method according to claim 28 wherein the compound of the general formula (I) and/or pharmaceutically acceptable salts thereof is administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M.

31. Method for detecting compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases comprising:
- 5 a) contacting a test compound with at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1; and
- 10 b) detecting the activity of said human cellular protein kinase, phosphatase or cellular signal transduction molecule.
32. Method for preventing and/or treating prion infections and/or diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which activates at least
- 15 partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or which activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the
- 20 group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
33. Method for regulating the production of prions in an individual comprising the step of administering an individual a pharmaceutically effective amount of at
- 25 least one pharmaceutically active agent wherein said agent activates at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or wherein said
- 30 agent at least partially activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 35 34. Method for regulating the production of prions in cells comprising the step of administering the cells a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent activates at least partially

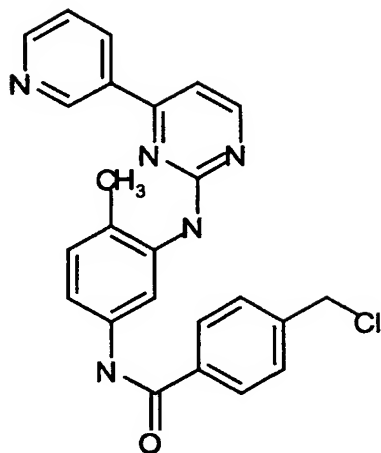
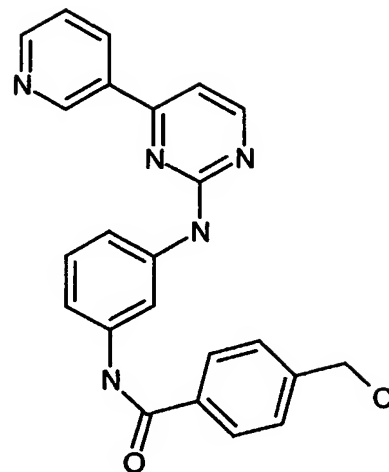
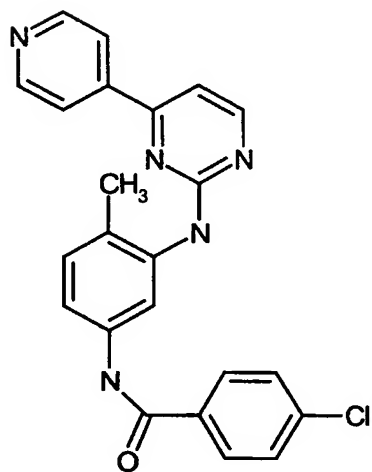
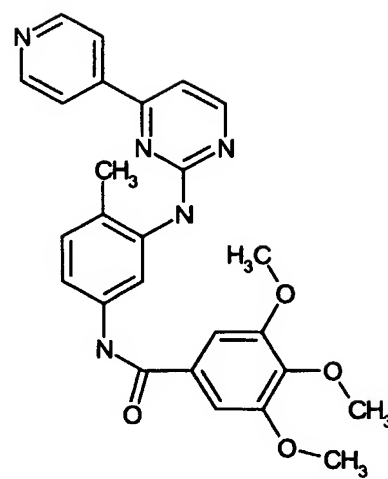
the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or wherein said agent at least partially activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in the cells.

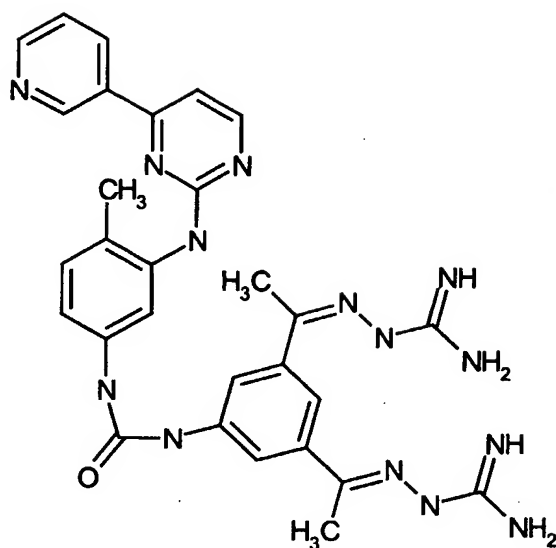
- 5
- 10 35. Method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in an individual comprising the step of administering the individual a
- 15 pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent inhibits at least partially the transcription of DNA or the translation of RNA.
- 20 36. Method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in the cells comprising the step of administering the cells a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent
- 25 inhibits at least partially the transcription of DNA or the translation of RNA.
- 30 37. Oligonucleotide that binds to the DNA or RNA encoding a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 35 38. Method according to claim 22, 23, 24, 25, 35 or 36 wherein the agent is a oligonucleotide which binds to the DNA and/or RNA encoding a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

39. Method according to claims 16, 17, 18, 19, 22, 24, 32, 33, or 35 wherein said individual is a human or ruminant.
- 5 40. Method according to any one of claims 17, 18, 19, 21, 22, 23, 31, or 32 wherein said prion infection and/or prion disease is selected from the group comprising Scrapie, TME, CWD, BSE, vCJD, CJD, GSS, FFI, Kuru, and Alpers Syndrome.
- 10 41. Method according to claim 40 wherein said prion infection and/or prion disease is BSE, vCJD, or CJD.
- 15 42. A solid support useful for detecting prion infections and/or diseases in an individual, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 20 43. A solid support useful for detecting prion infections and/or diseases in cells, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 25 44. A solid support useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized oligonucleotide, wherein said oligonucleotide encodes one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 30 45. A solid support useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the

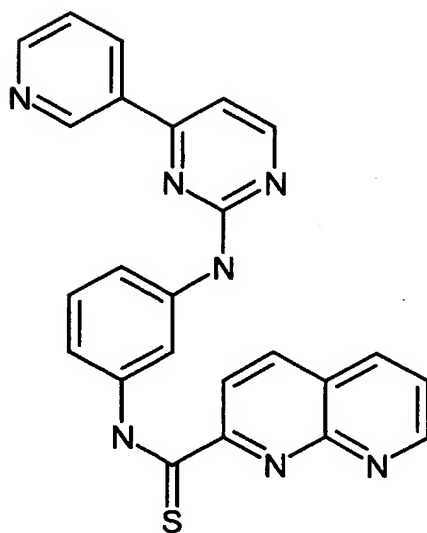
group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

- 5 46. Composition useful for the prophylaxis and/or treatment of an individual afflicted with prions comprising at least one agent capable of inhibiting at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 10 47. Composition useful for the prophylaxis and/or treatment of an individual afflicted with prions comprising at least one agent capable of activating or stimulating at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 15 48. Composition according claim 46 or 47, wherein the agent is at least one compound of the general formula (I) and/or pharmaceutically acceptable salts thereof.
- 20

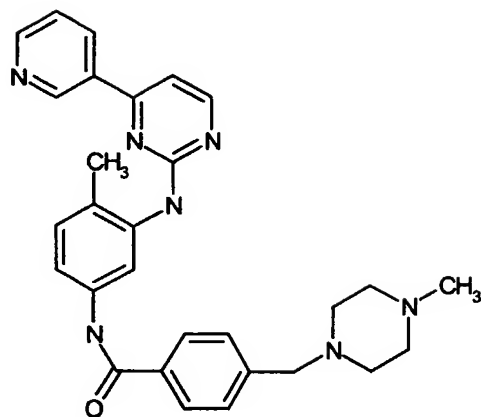
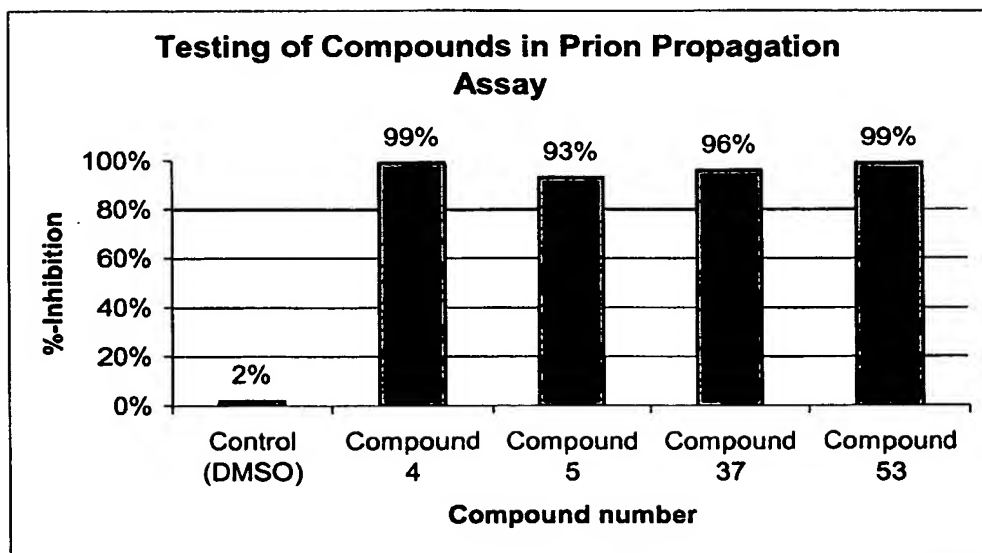
Fig. 1**Compound 4****Compound 5****Compound 37****Compound 52**



Compound 84



Compound 88

Fig. 2**Compound 53 (Gleevec™)****Fig. 3**

SEQUENCE LISTING

<110> Axxima Pharmaceuticals AG

5 <120> Human cellular protein kinases and phosphatases as
targets for diagnosis and treatment of prion diseases

<130> AXX-P11001-WO

10 <140> 0111858.5
<141> 2001-05-16

<150> US 60/293,528
<151> 2001-05-29

15 <160>

<170> PatentIn Ver. 2.1

20 <210> 1
<211> 2662
<212> DNA
<213> Homo sapiens

25 <220> FGF-R1
<223> Description of Sequence: N/A

<400> 1

tcagtttgaa aaggaggatc gagctcactc gtggagtatc catggagatg tggagccttg 60
30 tcaccaacct ctaactgcag aactgggatg tggagctgga agtgccctct cttctgggct 120
gtgctggtca cagccacact ctgcaccgct aggcctgccc cgaccttgcc tgaacaagcc 180
cagccctggg gagcccctgt ggaagtggag tccttctctgg tccaccccgg tgacctgctg 240
cagcttcgct gtcggctgcg ggacgatgtg cagagcatca actggctgcg ggacggggtg 300
cagctggcgg aaagcaaccg caccgcgcatc acaggggagg aggtggaggt gcaggactcc 360
35 gtgcccgcag actccggcct ctatgcttgc gtaaccagca gcccctcggg cagtgcacacc 420
acctacttct ccgtcaatgt ttcagatgct ctcccctcct cggaggatga tgatgatgat 480
gatgactcct cttcagagga gaaagaaaca gataacacca aaccaaaccg tatgcccgtg 540
gctccatatt ggacatcccc agaaaagatg gaaaagaaat tgcattgcagt gccggctgcc 600

aagacagtga agttcaaagt cccttccagt gggaccccaa accccacact gcgctgggtg 660
 aaaaatggca aagaattcaa acctgaccac agaattggag gctacaaggt ccgttatgcc 720
 acctggagca tcataatgga ctctgtggtg ccctctgaca agggcaacta cacctgcatt 780
 gtggagaatg agtacggcag catcaaccac acataccagc tggatgtcgt ggagcgggtc 840
 5 cctcaccgcc ccatcctgca agcagggttg cccgccaaca aaacagtggc cctgggtagc 900
 aacgtggagt tcatgtgtaa ggtgtacagt gaccgcagc cgcacatcca gtggctaaag 960
 cacatcgagg tgaatgggag caagattggc ccagacaacc tgccttatgt ccagatcctg 1020
 aagactgctg gagttaatac caccgacaaa gagatggagg tgcttcaactt aagaaatgtc 1080
 tcctttgagg acgcagggga gtatacgtgc ttggcgggta actctatcgg actctcccat 1140
 10 cactctgcat gggtgaccgt tctggaagcc ctggaagaga ggccggcagt gatgacctcg 1200
 cccctgtacc tggagatcat catctattgc acaggggcct tcctcatctc ctgcatggtg 1260
 gggtcgggtca tcgtctacaa gatgaagagt ggtaccaaga agagtgactt ccacagccag 1320
 atggctgtgc acaagctggc caagagcatc cctctgcgca gacaggtaac agtgtctgct 1380
 gactccagtg catccatgaa ctctggggtt cttctgggtc ggccatcacg gctctcctcc 1440
 15 agtgggactc ccatgctagc aggggtctct gagtatgagc ttcccgaaga ccctcgctgg 1500
 gagctgcctc gggacagact ggtcttaggc aaaccctgg gagagggctg ctttgggcag 1560
 gtggtgttgg cagaggctat cgggctggac aaggacaaac ccaaccgtgt gaccaaagt 1620
 gctgtgaaga tgttgaagtc ggacgcaaca gagaaagact tgtcagacct gatctcagaa 1680
 atggagatga tgaagatgat cgggaagcat aagaatatca tcaacctgct gggggcctgc 1740
 20 acgcaggatg gtcccttgta tgtcatcgtg gagtatgcct ccaagggcaa cctgcgggag 1800
 tacctgcagg cccggaggcc cccagggtcg gaatactgct acaaccccag ccacaacca 1860
 gaggagcagc tctcctccaa ggacctggtg tcctgcgcct accagggtggc ccgaggcatg 1920
 gagtatctgg cctccaagaa gtgcatacac cgagacctgg cagccaggaa tgtcctggtg 1980
 acagaggaca atgtgatgaa gatagcagac tttggcctcg cacgggacat tcaccacatc 2040
 25 gactactata aaaagacaac caacggccga ctgcctgtga agtggatggc acccgaggca 2100
 ttatttgacc ggatctacac ccaccagagt gatgtgtggt ctttcggggg gctcctgtgg 2160
 gagatcttca ctctgggcgg ctcccatac cccggtgtgc ctgtggagga acttttcaag 2220
 ctgctgaagg agggtcaccg catggacaag cccagtaact gcaccaacga gctgtacatg 2280
 atgatgcggg actgctggca tgcagtgcc tcacagagac ccaccttcaa gcagctggtg 2340
 30 gaagacctgg accgcatcgt ggccttgacc tccaaccagg agtacctgga cctgtccatg 2400
 cccctggacc agtactcccc cagctttccc gacacccgga gctctacgtg ctctcaggg 2460
 gaggattccg tcttctctca tgagccgctg cccgaggagc cctgcctgcc ccgacacca 2520
 gccagcttg ccaatggcgg actcaaacgc cgctgactgc caccacacg ccctcccag 2580
 actccaccgt cagctgtaac cctcaccac agcccctgcc tgggcccacc acctgtccgt 2640
 35 ccctgtcccc tttcctgctg gg 2662

<211> 822

<212> PRT

<213> Homo sapiens

5 <220>

<223> Description of Sequence: N/A

<400> 2

```

Met Trp Ser Trp Lys Cys Leu Leu Phe Trp Ala Val Leu Val Thr Ala
10      1              5              10              15
Thr Leu Cys Thr Ala Arg Pro Ser Pro Thr Leu Pro Glu Gln Ala Gln
      20              25              30
Pro Trp Gly Ala Pro Val Glu Val Glu Ser Phe Leu Val His Pro Gly
      35              40              45
15 Asp Leu Leu Gln Leu Arg Cys Arg Leu Arg Asp Asp Val Gln Ser Ile
      50              55              60
Asn Trp Leu Arg Asp Gly Val Gln Leu Ala Glu Ser Asn Arg Thr Arg
      65              70              75              80
Ile Thr Gly Glu Glu Val Glu Val Gln Asp Ser Val Pro Ala Asp Ser
20      85              90              95
Gly Leu Tyr Ala Cys Val Thr Ser Ser Pro Ser Gly Ser Asp Thr Thr
      100             105             110
Tyr Phe Ser Val Asn Val Ser Asp Ala Leu Pro Ser Ser Glu Asp Asp
      115             120             125
25 Asp Asp Asp Asp Asp Ser Ser Ser Glu Glu Lys Glu Thr Asp Asn Thr
      130             135             140
Lys Pro Asn Arg Met Pro Val Ala Pro Tyr Trp Thr Ser Pro Glu Lys
      145             150             155             160
Met Glu Lys Lys Leu His Ala Val Pro Ala Ala Lys Thr Val Lys Phe
30      165             170             175
Lys Cys Pro Ser Ser Gly Thr Pro Asn Pro Thr Leu Arg Trp Leu Lys
      180             185             190
Asn Gly Lys Glu Phe Lys Pro Asp His Arg Ile Gly Gly Tyr Lys Val
      195             200             205
35 Arg Tyr Ala Thr Trp Ser Ile Ile Met Asp Ser Val Val Pro Ser Asp
      210             215             220
Lys Gly Asn Tyr Thr Cys Ile Val Glu Asn Glu Tyr Gly Ser Ile Asn
      225             230             235             240

```

	His	Thr	Tyr	Gln	Leu	Asp	Val	Val	Glu	Arg	Ser	Pro	His	Arg	Pro	Ile	
					245					250					255		
	Leu	Gln	Ala	Gly	Leu	Pro	Ala	Asn	Lys	Thr	Val	Ala	Leu	Gly	Ser	Asn	
					260					265					270		
5	Val	Glu	Phe	Met	Cys	Lys	Val	Tyr	Ser	Asp	Pro	Gln	Pro	His	Ile	Gln	
					275					280					285		
	Trp	Leu	Lys	His	Ile	Glu	Val	Asn	Gly	Ser	Lys	Ile	Gly	Pro	Asp	Asn	
					290					295					300		
	Leu	Pro	Tyr	Val	Gln	Ile	Leu	Lys	Thr	Ala	Gly	Val	Asn	Thr	Thr	Asp	
10	305					310					315				320		
	Lys	Glu	Met	Glu	Val	Leu	His	Leu	Arg	Asn	Val	Ser	Phe	Glu	Asp	Ala	
					325					330					335		
	Gly	Glu	Tyr	Thr	Cys	Leu	Ala	Gly	Asn	Ser	Ile	Gly	Leu	Ser	His	His	
					340					345					350		
15	Ser	Ala	Trp	Leu	Thr	Val	Leu	Glu	Ala	Leu	Glu	Glu	Arg	Pro	Ala	Val	
					355					360					365		
	Met	Thr	Ser	Pro	Leu	Tyr	Leu	Glu	Ile	Ile	Ile	Tyr	Cys	Thr	Gly	Ala	
					370					375					380		
	Phe	Leu	Ile	Ser	Cys	Met	Val	Gly	Ser	Val	Ile	Val	Tyr	Lys	Met	Lys	
20	385					390					395				400		
	Ser	Gly	Thr	Lys	Lys	Ser	Asp	Phe	His	Ser	Gln	Met	Ala	Val	His	Lys	
					405					410					415		
	Leu	Ala	Lys	Ser	Ile	Pro	Leu	Arg	Arg	Gln	Val	Thr	Val	Ser	Ala	Asp	
					420					425					430		
25	Ser	Ser	Ala	Ser	Met	Asn	Ser	Gly	Val	Leu	Leu	Val	Arg	Pro	Ser	Arg	
					435					440					445		
	Leu	Ser	Ser	Ser	Gly	Thr	Pro	Met	Leu	Ala	Gly	Val	Ser	Glu	Tyr	Glu	
					450					455					460		
	Leu	Pro	Glu	Asp	Pro	Arg	Trp	Glu	Leu	Pro	Arg	Asp	Arg	Leu	Val	Leu	
30	465					470					475				480		
	Gly	Lys	Pro	Leu	Gly	Glu	Gly	Cys	Phe	Gly	Gln	Val	Val	Leu	Ala	Glu	
					485					490					495		
	Ala	Ile	Gly	Leu	Asp	Lys	Asp	Lys	Pro	Asn	Arg	Val	Thr	Lys	Val	Ala	
					500					505					510		
35	Val	Lys	Met	Leu	Lys	Ser	Asp	Ala	Thr	Glu	Lys	Asp	Leu	Ser	Asp	Leu	
					515					520					525		
	Ile	Ser	Glu	Met	Glu	Met	Met	Lys	Met	Ile	Gly	Lys	His	Lys	Asn	Ile	
					530					535					540		

Ile Asn Leu Leu Gly Ala Cys Thr Gln Asp Gly Pro Leu Tyr Val Ile
 545 550 555 560
 Val Glu Tyr Ala Ser Lys Gly Asn Leu Arg Glu Tyr Leu Gln Ala Arg
 565 570 575
 5 Arg Pro Pro Gly Leu Glu Tyr Cys Tyr Asn Pro Ser His Asn Pro Glu
 580 585 590
 Glu Gln Leu Ser Ser Lys Asp Leu Val Ser Cys Ala Tyr Gln Val Ala
 595 600 605
 Arg Gly Met Glu Tyr Leu Ala Ser Lys Lys Cys Ile His Arg Asp Leu
 10 610 615 620
 Ala Ala Arg Asn Val Leu Val Thr Glu Asp Asn Val Met Lys Ile Ala
 625 630 635 640
 Asp Phe Gly Leu Ala Arg Asp Ile His His Ile Asp Tyr Tyr Lys Lys
 645 650 655
 15 Thr Thr Asn Gly Arg Leu Pro Val Lys Trp Met Ala Pro Glu Ala Leu
 660 665 670
 Phe Asp Arg Ile Tyr Thr His Gln Ser Asp Val Trp Ser Phe Gly Val
 675 680 685
 Leu Leu Trp Glu Ile Phe Thr Leu Gly Gly Ser Pro Tyr Pro Gly Val
 20 690 695 700
 Pro Val Glu Glu Leu Phe Lys Leu Leu Lys Glu Gly His Arg Met Asp
 705 710 715 720
 Lys Pro Ser Asn Cys Thr Asn Glu Leu Tyr Met Met Met Arg Asp Cys
 725 730 735
 25 Trp His Ala Val Pro Ser Gln Arg Pro Thr Phe Lys Gln Leu Val Glu
 740 745 750
 Asp Leu Asp Arg Ile Val Ala Leu Thr Ser Asn Gln Glu Tyr Leu Asp
 755 760 765
 Leu Ser Met Pro Leu Asp Gln Tyr Ser Pro Ser Phe Pro Asp Thr Arg
 30 770 775 780
 Ser Ser Thr Cys Ser Ser Gly Glu Asp Ser Val Phe Ser His Glu Pro
 785 790 795 800
 Leu Pro Glu Glu Pro Cys Leu Pro Arg His Pro Ala Gln Leu Ala Asn
 805 810 815
 35 Gly Gly Leu Lys Arg Arg
 820

<210> 3

<211> 3840

<212> DNA

5 <213> Homo sapiens

<220> Abl

<223> Description of Sequence: N/A

10 <400> 3

ggccttcccc ctgcgaggat cgccgttggc ccgggttggc tttggaaagc ggcggtggct 60
ttgggcccggg ctccggcctcg ggaacgccag gggcccctgg gtgcggacgg gcgcggccag 120
gaggggggta aggcgcaggc ggcggcgggg cggggggcggg cctggcgggc gccctctccg 180
ggccctttgt taacaggcgc gtcccgcca gcggagacgc ggccgcctg ggcgggcgcg 240
15 ggcggcgggc ggcggtgagg gcggcctcg ggcggcgcc cggggggccg gccgagccgg 300
gcctgagccg ggcccgacc gagctgggag aggggctccg gcccgatcgt tcgcttggcg 360
caaaatgttg gagatctgcc tgaagctggg gggctgcaa tccaagaagg ggctgtcctc 420
gtcctccagc tgttatctgg aagaagccct tcagcggcca gtagcatctg actttgagcc 480
tcagggtctg agtgaagccg ctcggtgaa ctccaaggaa aaccttctcg ctggaccag 540
20 tgaaaatgac cccaaccttt tcgttgact gtatgatttt gtggccagtg gagataacac 600
tctaagcata actaaagggtg aaaagctccg ggtcttaggc tataatcaca atggggaatg 660
gtgtgaagcc caaaccaaaa atggccaagg ctgggtccca agcaactaca tcacgccagt 720
caacagtctg gagaaacact cctggtacca tgggcctgtg tcccgcaatg ccgctgagta 780
tccgctgagc agcgggatca atggcagctt cttggtgcgt gagagtgaga gcagtcctag 840
25 ccagaggtcc atctcgctga gatacgaagg gaggggtgtac cattacagga tcaacactgc 900
ttctgatggc aagctctacg tctcctccga gagccgcttc aacaccctgg ccgagttggg 960
tcatcatcat tcaacggtgg ccgacgggct catcaccacg ctccattatc cagcccaaaa 1020
gcgcaacaag cccactgtct atggtgtgtc cccaactac gacaagtggg agatggaacg 1080
cacggacatc accatgaagc acaagctggg cggggggccag tacggggagg tgtacgaggg 1140
30 cgtgtggaag aaatacagcc tgacggtggc cgtgaagacc ttgaaggagg acaccatgga 1200
ggtggaagag ttcttgaaag aagctgcagt catgaaagag atcaaacacc ctaacctagt 1260
gcagctcctt ggggtctgca cccgggagcc cccgttctat atcatcactg agttcatgac 1320
ctacgggaac ctcttgact acctgaggga gtgcaaccgg caggaggtga acgccgtggg 1380
gctgctgtac atggccactc agatctcgtc agccatggag tacctagaga agaaaaactt 1440
35 catccacaga gatcttgctg cccgaaactg cctggtaggg gagaaccact tgggtgaagg 1500
agctgatttt ggcctgagca ggttgatgac aggggacacc tacacagccc atgctggagc 1560
caagttcccc atcaaagga ctgcaccga gagcctggcc tacaacaagt tctccatcaa 1620
gtccgacgtc tgggcatttg gagtattgct ttgggaaatt gctacctatg gcatgtcccc 1680

ttacccggga attgaccgtt cccaggtgta tgagctgcta gagaaggact accgcatgaa 1740
gcgcccagaa ggctgcccag agaaggtcta tgaactcatg cgagcatggt ggcagtggaa 1800
tccctctgac cggccctcct ttgctgaaat ccaccaagcc tttgaaacaa tggtccagga 1860
atccagtatc tcagacgaag tggaaaagga gctggggaaa caaggcgtcc gtggggctgt 1920
5 gactaccttg ctgcaggccc cagagctgcc caccaagacg aggacctcca ggagagctgc 1980
agagcacaga gacaccactg acgtgcctga gatgcctcac tccaagggcc agggagagag 2040
cgatcctctg gaccatgagc ctgccgtgtc tccattgtct cctcgaaaag agcgagggtcc 2100
cccgaggggc ggctgaatg aagatgagcg ctttctcccc aaagacaaaa agaccaactt 2160
gttcagcgcc ttgatcaaga agaagaagaa gacagcccca accctccca aacgcagcag 2220
10 ctcttcccg gagatggacg gccagccgga gcgcagaggg gccggcgagg aagagggccg 2280
agacatcagc aacggggcac tggctttcac ccccttgac acagctgacc cagccaagt 2340
cccaaagccc agcaatgggg ctgggggtccc caatggagcc ctccgggagt ccgggggctc 2400
aggcttcccg tctccccacc tgtggaagaa gtccagcacg ctgaccagca gccgcctagc 2460
caccggcgag gaggagggcg gtggcagctc cagcaagcgc ttctgcgct cttgtccgt 2520
15 ctctgcggt ccccatgggg ccaaggacac ggagtggagg tcagtcacgc tgcctcgga 2580
cttgagctcc acgggaagac agtttgactc gtccacattt ggagggcaca aaagtgagaa 2640
gccggctctg cctcggaaga gggcagggga gaacaggtct gaccaggtga cccgaggcac 2700
agtaacgcct ccccccaggc tggtgaaaaa gaatgaggaa gctgctgatg aggtcttcaa 2760
agacatcatg gagtccagcc cgggctccag cccgcccaac ctgactcaa aaccctccg 2820
20 gcggcaggtc accgtggccc ctgcctcggg cctccccac aaggaagaag cctggaaagg 2880
cagtgcctta gggaccctg ctgcagctga gccagtgacc cccaccagca aagcaggctc 2940
aggtgcacca aggggcacca gcaaggccc cgccgaggag tccagagtga ggaggcaca 3000
gcactcctct gagtgcgag ggagggacaa ggggaaattt tccaagctca aacctgccc 3060
gccgccccca ccagcagcct ctgcaggga ggctggagga aagccctcgc agaggcccg 3120
25 ccaggaggct gccggggagg cagtcttggg cgcaaagaca aaagccacga gtctggttga 3180
tgctgtgaac agtgacgctg ccaagcccag ccagccggca gagggcctca aaaagcccgt 3240
gctcccggcc actccaaagc cacacccgc caagccgtcg gggacccca tcagcccagc 3300
ccccgttccc ctttccacgt tgccatcagc atcctcggcc ttggcagggg accagccgtc 3360
ttccactgcc ttcacccctc tcatatcaac ccgagtgtct cttcggaaaa cccgccagcc 3420
30 tccagagcgg gccagcggcg ccacaccaa gggcgtgggtc ttggacagca ccgaggcgct 3480
gtgcctcgcc atctctggga actccgagca gatggccagc cacagcgagc tgctggaggc 3540
cggcaaaaac ctctacacgt tctgcgtgag ctatgtggat tccatccagc aaatgaggaa 3600
caagtttgcc ttccgagagg ccacacaaa actggagaat aatctccggg agcttcagat 3660
ctgcccggcg tcagcaggca gtggtccggc ggccactcag gacttcagca agctcctcag 3720
35 ttcggtgaag gaaatcagtg acatagtga gaggtagcag cagtcagggg tcaggtgtca 3780
ggcccgctcg agctgcctgc agcacatgcg ggctcgccca taccatgac agtggctgag 3840

<210> 4

<211> 1130

<212> PRT

<213> Homo sapiens

5

<220>

<223> Description of Sequence: N/A

<400> 4

10 Met Leu Glu Ile Cys Leu Lys Leu Val Gly Cys Lys Ser Lys Lys Gly
 1 5 10 15
 Leu Ser Ser Ser Ser Ser Cys Tyr Leu Glu Glu Ala Leu Gln Arg Pro
 20 25 30
 Val Ala Ser Asp Phe Glu Pro Gln Gly Leu Ser Glu Ala Ala Arg Trp
 15 35 40 45
 Asn Ser Lys Glu Asn Leu Leu Ala Gly Pro Ser Glu Asn Asp Pro Asn
 50 55 60
 Leu Phe Val Ala Leu Tyr Asp Phe Val Ala Ser Gly Asp Asn Thr Leu
 65 70 75 80
 20 Ser Ile Thr Lys Gly Glu Lys Leu Arg Val Leu Gly Tyr Asn His Asn
 85 90 95
 Gly Glu Trp Cys Glu Ala Gln Thr Lys Asn Gly Gln Gly Trp Val Pro
 100 105 110
 Ser Asn Tyr Ile Thr Pro Val Asn Ser Leu Glu Lys His Ser Trp Tyr
 25 115 120 125
 His Gly Pro Val Ser Arg Asn Ala Ala Glu Tyr Pro Leu Ser Ser Gly
 130 135 140
 Ile Asn Gly Ser Phe Leu Val Arg Glu Ser Glu Ser Ser Pro Ser Gln
 145 150 155 160
 30 Arg Ser Ile Ser Leu Arg Tyr Glu Gly Arg Val Tyr His Tyr Arg Ile
 165 170 175
 Asn Thr Ala Ser Asp Gly Lys Leu Tyr Val Ser Ser Glu Ser Arg Phe
 180 185 190
 Asn Thr Leu Ala Glu Leu Val His His His Ser Thr Val Ala Asp Gly
 35 195 200 205
 Leu Ile Thr Thr Leu His Tyr Pro Ala Pro Lys Arg Asn Lys Pro Thr
 210 215 220
 Val Tyr Gly Val Ser Pro Asn Tyr Asp Lys Trp Glu Met Glu Arg Thr

	225					230						235				240
	Asp	Ile	Thr	Met	Lys	His	Lys	Leu	Gly	Gly	Gly	Gln	Tyr	Gly	Glu	Val
					245					250					255	
	Tyr	Glu	Gly	Val	Trp	Lys	Lys	Tyr	Ser	Leu	Thr	Val	Ala	Val	Lys	Thr
5				260					265					270		
	Leu	Lys	Glu	Asp	Thr	Met	Glu	Val	Glu	Glu	Phe	Leu	Lys	Glu	Ala	Ala
				275					280					285		
	Val	Met	Lys	Glu	Ile	Lys	His	Pro	Asn	Leu	Val	Gln	Leu	Leu	Gly	Val
				290				295				300				
10	Cys	Thr	Arg	Glu	Pro	Pro	Phe	Tyr	Ile	Ile	Thr	Glu	Phe	Met	Thr	Tyr
	305					310					315					320
	Gly	Asn	Leu	Leu	Asp	Tyr	Leu	Arg	Glu	Cys	Asn	Arg	Gln	Glu	Val	Asn
					325					330					335	
	Ala	Val	Val	Leu	Leu	Tyr	Met	Ala	Thr	Gln	Ile	Ser	Ser	Ala	Met	Glu
15				340					345					350		
	Tyr	Leu	Glu	Lys	Lys	Asn	Phe	Ile	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn
				355				360					365			
	Cys	Leu	Val	Gly	Glu	Asn	His	Leu	Val	Lys	Val	Ala	Asp	Phe	Gly	Leu
				370			375					380				
20	Ser	Arg	Leu	Met	Thr	Gly	Asp	Thr	Tyr	Thr	Ala	His	Ala	Gly	Ala	Lys
	385					390					395					400
	Phe	Pro	Ile	Lys	Trp	Thr	Ala	Pro	Glu	Ser	Leu	Ala	Tyr	Asn	Lys	Phe
					405					410					415	
	Ser	Ile	Lys	Ser	Asp	Val	Trp	Ala	Phe	Gly	Val	Leu	Leu	Trp	Glu	Ile
25				420					425					430		
	Ala	Thr	Tyr	Gly	Met	Ser	Pro	Tyr	Pro	Gly	Ile	Asp	Arg	Ser	Gln	Val
				435				440					445			
	Tyr	Glu	Leu	Leu	Glu	Lys	Asp	Tyr	Arg	Met	Lys	Arg	Pro	Glu	Gly	Cys
				450			455					460				
30	Pro	Glu	Lys	Val	Tyr	Glu	Leu	Met	Arg	Ala	Cys	Trp	Gln	Trp	Asn	Pro
	465					470					475					480
	Ser	Asp	Arg	Pro	Ser	Phe	Ala	Glu	Ile	His	Gln	Ala	Phe	Glu	Thr	Met
					485					490					495	
	Phe	Gln	Glu	Ser	Ser	Ile	Ser	Asp	Glu	Val	Glu	Lys	Glu	Leu	Gly	Lys
35				500					505					510		
	Gln	Gly	Val	Arg	Gly	Ala	Val	Thr	Thr	Leu	Leu	Gln	Ala	Pro	Glu	Leu
				515				520					525			
	Pro	Thr	Lys	Thr	Arg	Thr	Ser	Arg	Arg	Ala	Ala	Glu	His	Arg	Asp	Thr

	530					535						540				
	Thr	Asp	Val	Pro	Glu	Met	Pro	His	Ser	Lys	Gly	Gln	Gly	Glu	Ser	Asp
	545					550					555					560
5	Pro	Leu	Asp	His	Glu	Pro	Ala	Val	Ser	Pro	Leu	Leu	Pro	Arg	Lys	Glu
					565					570					575	
	Arg	Gly	Pro	Pro	Glu	Gly	Gly	Leu	Asn	Glu	Asp	Glu	Arg	Leu	Leu	Pro
					580				585					590		
	Lys	Asp	Lys	Lys	Thr	Asn	Leu	Phe	Ser	Ala	Leu	Ile	Lys	Lys	Lys	Lys
		595						600					605			
10	Lys	Thr	Ala	Pro	Thr	Pro	Pro	Lys	Arg	Ser	Ser	Ser	Phe	Arg	Glu	Met
	610						615					620				
	Asp	Gly	Gln	Pro	Glu	Arg	Arg	Gly	Ala	Gly	Glu	Glu	Glu	Gly	Arg	Asp
	625					630					635					640
	Ile	Ser	Asn	Gly	Ala	Leu	Ala	Phe	Thr	Pro	Leu	Asp	Thr	Ala	Asp	Pro
15					645					650					655	
	Ala	Lys	Ser	Pro	Lys	Pro	Ser	Asn	Gly	Ala	Gly	Val	Pro	Asn	Gly	Ala
					660				665					670		
	Leu	Arg	Glu	Ser	Gly	Gly	Ser	Gly	Phe	Arg	Ser	Pro	His	Leu	Trp	Lys
		675						680					685			
20	Lys	Ser	Ser	Thr	Leu	Thr	Ser	Ser	Arg	Leu	Ala	Thr	Gly	Glu	Glu	Glu
	690						695					700				
	Gly	Gly	Gly	Ser	Ser	Ser	Lys	Arg	Phe	Leu	Arg	Ser	Cys	Ser	Val	Ser
	705					710					715					720
	Cys	Val	Pro	His	Gly	Ala	Lys	Asp	Thr	Glu	Trp	Arg	Ser	Val	Thr	Leu
25					725					730					735	
	Pro	Arg	Asp	Leu	Gln	Ser	Thr	Gly	Arg	Gln	Phe	Asp	Ser	Ser	Thr	Phe
					740				745					750		
	Gly	Gly	His	Lys	Ser	Glu	Lys	Pro	Ala	Leu	Pro	Arg	Lys	Arg	Ala	Gly
		755						760					765			
30	Glu	Asn	Arg	Ser	Asp	Gln	Val	Thr	Arg	Gly	Thr	Val	Thr	Pro	Pro	Pro
	770						775					780				
	Arg	Leu	Val	Lys	Lys	Asn	Glu	Glu	Ala	Ala	Asp	Glu	Val	Phe	Lys	Asp
	785					790					795					800
	Ile	Met	Glu	Ser	Ser	Pro	Gly	Ser	Ser	Pro	Pro	Asn	Leu	Thr	Pro	Lys
35					805					810					815	
	Pro	Leu	Arg	Arg	Gln	Val	Thr	Val	Ala	Pro	Ala	Ser	Gly	Leu	Pro	His
					820				825					830		
	Lys	Glu	Glu	Ala	Trp	Lys	Gly	Ser	Ala	Leu	Gly	Thr	Pro	Ala	Ala	Ala

	835	840	845
	Glu Pro Val Thr Pro Thr Ser Lys Ala Gly Ser Gly Ala Pro Arg Gly		
	850	855	860
	Thr Ser Lys Gly Pro Ala Glu Glu Ser Arg Val Arg Arg His Lys His		
5	865	870	875 880
	Ser Ser Glu Ser Pro Gly Arg Asp Lys Gly Lys Leu Ser Lys Leu Lys		
	885	890	895
	Pro Ala Pro Pro Pro Pro Pro Ala Ala Ser Ala Gly Lys Ala Gly Gly		
	900	905	910
10	Lys Pro Ser Gln Arg Pro Gly Gln Glu Ala Ala Gly Glu Ala Val Leu		
	915	920	925
	Gly Ala Lys Thr Lys Ala Thr Ser Leu Val Asp Ala Val Asn Ser Asp		
	930	935	940
	Ala Ala Lys Pro Ser Gln Pro Ala Glu Gly Leu Lys Lys Pro Val Leu		
15	945	950	955 960
	Pro Ala Thr Pro Lys Pro His Pro Ala Lys Pro Ser Gly Thr Pro Ile		
	965	970	975
	Ser Pro Ala Pro Val Pro Leu Ser Thr Leu Pro Ser Ala Ser Ser Ala		
	980	985	990
20	Leu Ala Gly Asp Gln Pro Ser Ser Thr Ala Phe Ile Pro Leu Ile Ser		
	995	1000	1005
	Thr Arg Val Ser Leu Arg Lys Thr Arg Gln Pro Pro Glu Arg Ala Ser		
	1010	1015	1020
	Gly Ala Ile Thr Lys Gly Val Val Leu Asp Ser Thr Glu Ala Leu Cys		
25	1025	1030	1035 1040
	Leu Ala Ile Ser Gly Asn Ser Glu Gln Met Ala Ser His Ser Ala Val		
	1045	1050	1055
	Leu Glu Ala Gly Lys Asn Leu Tyr Thr Phe Cys Val Ser Tyr Val Asp		
	1060	1065	1070
30	Ser Ile Gln Gln Met Arg Asn Lys Phe Ala Phe Arg Glu Ala Ile Asn		
	1075	1080	1085
	Lys Leu Glu Asn Asn Leu Arg Glu Leu Gln Ile Cys Pro Ala Ser Ala		
	1090	1095	1100
	Gly Ser Gly Pro Ala Ala Thr Gln Asp Phe Ser Lys Leu Leu Ser Ser		
35	1105	1110	1115 1120
	Val Lys Glu Ile Ser Asp Ile Val Gln Arg		
	1125	1130	

<210> 5

<211> 1461

5 <212> DNA

<213> Homo sapiens

<220> MKK7

<223> Description of Sequence: N/A

10

<400> 5

15

20

25

30

35

```

aggcgggtgtt tgtctgccgg actgacgggc ggccggggcgg tgcgcggcgg cggtggcggc 60
ggggaaaatg gcggcgtcct ccctggaaca gaagctgtcc cgcctggaag caaagctgaa 120
gcaggagaac cgggaggccc ggcggaggat cgacctcaac ctggatatca gccccagcg 180
gcccaggccc accctgcagc tcccgtggc caacgatggg ggcagccgct cgccatcctc 240
agagagctcc ccgcagcacc ccacgcccc cgccggccc cgccacatgc tggggctccc 300
gtcaaccctg ttcacacccc gcagcatgga gagcattgag attgaccaga agctgcagga 360
gatcatgaag cagacgggct acctgaccat cggggggccag cgctaccagg cagaaatcaa 420
cgacctggag aacttgggcg agatgggcag cggcacctgc ggccagggtg ggaagatgcg 480
cttcgggaag accggccacg tcattgccgt taagcaaatg cggcgctccg ggaacaagga 540
ggagaacaag cgcatcctca tggacctgga tgtggtgctg aagagccacg actgccccta 600
catcgtgcag tgctttggga cgttcatcac caacacggac gtcttcatcg ccatggagct 660
catgggcacc tgcgctgaga agctcaagaa gcggatgcag ggccccatcc ccgagcgcat 720
tctgggcaag atgacagtgg cgattgtgaa ggcgctgtac tacctgaagg agaagcacgg 780
tgtcatccac cgcgacgtca agccctccaa catcctgctg gacgagcggg gccagatcaa 840
gttctgcgac ttcggcatca gcggccgcct ggtggactcc aaagccaaga cgcggagcgc 900
cggctgtgcc gcctacatgg cacccgagcg cattgacccc ccagacccca ccaagccgga 960
ctatgacatc cgggccgacg tatggagcct gggcatctcg ctggtggagc tggcaacagg 1020
acagtttccc tacaagaact gcaagacgga ctttgaggtc ctcaccaaag tcctacagga 1080
agagcccccg cttctgcccg gacacatggg cttctcgggg gacttccagt ccttcgtcaa 1140
agactgcctt actaaagatc acaggaagag accaaagtat aataagctac ttgaacacag 1200
cttcatcaag cgctacgaga cgctggaggt ggacgtggcg tcttggttca aggatgtcat 1260
ggcgaagact gagtcaccgc ggactagcgg cgtcctgagc cagccccacc tgcccttctt 1320
caggtagctg cttggcgggc gccagcccca cagggggcca ggggcatggc cacaggcccc 1380
cctccccact tggccacca gctgcctgcc aggggagacc tgggacctgg acggccacct 1440
aggactgagg acagagagtg g

```

1461

<210> 6

<211> 419

<212> PRT

<213> Homo sapiens

5

<220>

<223> Description of Sequence: N/A

<400> 6

10 Met Ala Ala Ser Ser Leu Glu Gln Lys Leu Ser Arg Leu Glu Ala Lys
 1 5 10 15
 Leu Lys Gln Glu Asn Arg Glu Ala Arg Arg Arg Ile Asp Leu Asn Leu
 20 25 30
 Asp Ile Ser Pro Gln Arg Pro Arg Pro Thr Leu Gln Leu Pro Leu Ala
 15 35 40 45
 Asn Asp Gly Gly Ser Arg Ser Pro Ser Ser Glu Ser Ser Pro Gln His
 50 55 60
 Pro Thr Pro Pro Ala Arg Pro Arg His Met Leu Gly Leu Pro Ser Thr
 65 70 75 80
 20 Leu Phe Thr Pro Arg Ser Met Glu Ser Ile Glu Ile Asp Gln Lys Leu
 85 90 95
 Gln Glu Ile Met Lys Gln Thr Gly Tyr Leu Thr Ile Gly Gly Gln Arg
 100 105 110
 Tyr Gln Ala Glu Ile Asn Asp Leu Glu Asn Leu Gly Glu Met Gly Ser
 25 115 120 125
 Gly Thr Cys Gly Gln Val Trp Lys Met Arg Phe Arg Lys Thr Gly His
 130 135 140
 Val Ile Ala Val Lys Gln Met Arg Arg Ser Gly Asn Lys Glu Glu Asn
 145 150 155 160
 30 Lys Arg Ile Leu Met Asp Leu Asp Val Val Leu Lys Ser His Asp Cys
 165 170 175
 Pro Tyr Ile Val Gln Cys Phe Gly Thr Phe Ile Thr Asn Thr Asp Val
 180 185 190
 Phe Ile Ala Met Glu Leu Met Gly Thr Cys Ala Glu Lys Leu Lys Lys
 35 195 200 205
 Arg Met Gln Gly Pro Ile Pro Glu Arg Ile Leu Gly Lys Met Thr Val
 210 215 220
 Ala Ile Val Lys Ala Leu Tyr Tyr Leu Lys Glu Lys His Gly Val Ile

```

225          230          235          240
His Arg Asp Val Lys Pro Ser Asn Ile Leu Leu Asp Glu Arg Gly Gln
          245          250          255
Ile Lys Phe Cys Asp Phe Gly Ile Ser Gly Arg Leu Val Asp Ser Lys
5          260          265          270
Ala Lys Thr Arg Ser Ala Gly Cys Ala Ala Tyr Met Ala Pro Glu Arg
          275          280          285
Ile Asp Pro Pro Asp Pro Thr Lys Pro Asp Tyr Asp Ile Arg Ala Asp
          290          295          300
10 Val Trp Ser Leu Gly Ile Ser Leu Val Glu Leu Ala Thr Gly Gln Phe
305          310          315          320
Pro Tyr Lys Asn Cys Lys Thr Asp Phe Glu Val Leu Thr Lys Val Leu
          325          330          335
Gln Glu Glu Pro Pro Leu Leu Pro Gly His Met Gly Phe Ser Gly Asp
15          340          345          350
Phe Gln Ser Phe Val Lys Asp Cys Leu Thr Lys Asp His Arg Lys Arg
          355          360          365
Pro Lys Tyr Asn Lys Leu Leu Glu His Ser Phe Ile Lys Arg Tyr Glu
          370          375          380
20 Thr Leu Glu Val Asp Val Ala Ser Trp Phe Lys Asp Val Met Ala Lys
385          390          395          400
Thr Glu Ser Pro Arg Thr Ser Gly Val Leu Ser Gln Pro His Leu Pro
          405          410          415
Phe Phe Arg

25

<210> 7
<211> 1050
30 <212> DNA
<213> Homo sapiens

<220> CDC2
<223> Description of Sequence: N/A
35
<400> 7
ggggggggggg ggcacttggc ttcaaagctg gctcttggaa attgagcggg gacgagcggc 60
ttgttqtatg tgcctgtcgg ccgccgcgga ataataagcc gggatctacc ataccattga 120

```

ctaactatgg aagattatac caaaatagag aaaattggag aaggtagccta tggagttgtg 180
tataagggtta gacacaaaac tacaggtcaa gtggttagcca tgaaaaaat cagactagaa 240
agtgaagagg aaggggttcc tagtactgca attcgggaaa tttctctatt aaaggaactt 300
cgtcatccaa atatagtcag tcttcaggat gtgcttatgc aggattccag gttatatctc 360
5 atctttgagt ttctttccat ggatctgaag aaatacttgg attctatccc tcctggtcag 420
tacatggatt cttcacttgt taagagttat ttataccaaa tcctacaggg gattgtgttt 480
tgtcactcta gaagagttct tcacagagac ttaaaacctc aaaatctctt gattgatgac 540
aaaggaacaa ttaaactggc tgattttggc cttgccagag cttttggaat acctatcaga 600
gtatatacac atgaggtagt aacactctgg tacagatctc cagaagtatt gctgggggtca 660
10 gctcgttact caactccagt tgacatttgg agtataggca ccatatttgc tgaactagca 720
actaagaaac cacttttcca tggggattca gaaattgatc aactcttcag gattttcaga 780
gctttgggca ctccaataa tgaagtgtgg ccagaagtgg aatctttaca ggactataag 840
aatacatttc ccaaattgaa accaggaagc ctagcatccc atgtcaaaaa cttggatgaa 900
aatggcttgg atttgctctc gaaaatgtta atctatgatc cagccaaacg aatttctggc 960
15 aaaatggcac tgaatcatcc atattttaat gatttggaca atcagattaa gaagatgtag 1020
ctttctgaca aaaagtttcc atatgttatg 1050

<210> 8

20 <211> 297

<212> PRT

<213> Homo sapiens

<220>

25 <223> Description of Sequence: N/A

<400> 8

Met Glu Asp Tyr Thr Lys Ile Glu Lys Ile Gly Glu Gly Thr Tyr Gly
1 5 10 15
30 Val Val Tyr Lys Gly Arg His Lys Thr Thr Gly Gln Val Val Ala Met
20 25 30
Lys Lys Ile Arg Leu Glu Ser Glu Glu Gly Val Pro Ser Thr Ala
35 40 45
Ile Arg Glu Ile Ser Leu Leu Lys Glu Leu Arg His Pro Asn Ile Val
35 50 55 60
Ser Leu Gln Asp Val Leu Met Gln Asp Ser Arg Leu Tyr Leu Ile Phe
65 70 75 80
Glu Phe Leu Ser Met Asp Leu Lys Lys Tyr Leu Asp Ser Ile Pro Pro

85 90 95
 Gly Gln Tyr Met Asp Ser Ser Leu Val Lys Ser Tyr Leu Tyr Gln Ile
 100 105 110
 Leu Gln Gly Ile Val Phe Cys His Ser Arg Arg Val Leu His Arg Asp
 5 115 120 125
 Leu Lys Pro Gln Asn Leu Leu Ile Asp Asp Lys Gly Thr Ile Lys Leu
 130 135 140
 Ala Asp Phe Gly Leu Ala Arg Ala Phe Gly Ile Pro Ile Arg Val Tyr
 145 150 155 160
 10 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ser Pro Glu Val Leu Leu
 165 170 175
 Gly Ser Ala Arg Tyr Ser Thr Pro Val Asp Ile Trp Ser Ile Gly Thr
 180 185 190
 Ile Phe Ala Glu Leu Ala Thr Lys Lys Pro Leu Phe His Gly Asp Ser
 15 195 200 205
 Glu Ile Asp Gln Leu Phe Arg Ile Phe Arg Ala Leu Gly Thr Pro Asn
 210 215 220
 Asn Glu Val Trp Pro Glu Val Glu Ser Leu Gln Asp Tyr Lys Asn Thr
 225 230 235 240
 20 Phe Pro Lys Trp Lys Pro Gly Ser Leu Ala Ser His Val Lys Asn Leu
 245 250 255
 Asp Glu Asn Gly Leu Asp Leu Leu Ser Lys Met Leu Ile Tyr Asp Pro
 260 265 270
 Ala Lys Arg Ile Ser Gly Lys Met Ala Leu Asn His Pro Tyr Phe Asn
 25 275 280 285
 Asp Leu Asp Asn Gln Ile Lys Lys Met
 290 295

30

<210> 9

<211> 1480

<212> DNA

<213> Homo sapiens

35

<220> CamKI

<223> Description of Sequence: N/A

<400> 9

5 gaattccgag caagagcgcg ggcgggtggc ccaggcacgc agcgggtgag gaccgcgccc 60
 acagctcggc gccaaaccacc gcgggcctcc cagccagccc cgcnnngagc cgcaggancc 120
 ctggctgtgg tcggggggca gtgggccatg ctgggggcag tggaaagccc caggtggaag 180
 10 caggcggagg acattagaga catctacgac ttccgagatg ttctgggcac gggggccttc 240
 tcggaggtga tcctggcaga agataagagg acgcagaagc tgggtggccat caaatgcatt 300
 gccaaaggagg ccctggaggg caaggaaggc agcatggaga atgagattgc tgtcctgcac 360
 aagatcaagc accccaacat tgtagccctg gatgacatct atgagagtgg gggccacctc 420
 tacctcatca tgcagctggt gtcgggtggg gagctctttg accgtattgt ggaaaaaggc 480
 15 ttctacacgg agcgggacgc cagccgcctc atcttccagg tgctggatgc tgtgaaatac 540
 ctgcatgacc tgggcattgt acaccgggat ctcaagccag agaatctgct gtactacagc 600
 ctggatgaag actccaaaat catgatctcc gactttggcc tctccaagat ggaggacccg 660
 ggcagtgtgc tctccaccgc ctgtggaact ccgggatacg tggcccctga agtcctggcc 720
 cagaagccct acagcaaggc tgtggattgc tgggccatag gtgtcatcgc ctacatcttg 780
 20 ctctgcggtt accctccctt ctatgacgag aatgatgcca aactctttga acagattttg 840
 aaggccgagt acgagtttga ctctccttac tgggacgaca tctctgactc tgccaaagat 900
 ttcacccggc acttgatgga gaaggacca gagaaaagat tcacctgtga gcaggccttg 960
 cagcacccat ggattgcagg agatacagct ctagataaga atatccacca gtcggtgagt 1020
 gagcagatca agaagaactt tgccaagagc aagtggaagc aagccttcaa tgccacggct 1080
 25 gtggtgcggc acatgaggaa actgcagctg ggcaccagcc aggaggggca ggggcagacg 1140
 gcgagccatg gggagctgct gacaccagtg gctggggggc cggcagctgg ctgttgctgt 1200
 cgagactgct gcgtggagcc gggcacagaa ctgtcccca cactgccccca ccagctctag 1260
 ggccctggac ctcgggtcat gatcctctgc gtgggagggc ttggggggcca gcctgctccc 1320
 ctccctccc tgaaccggga gtttctctgc cctgtccct cctcacctgc ttccctacca 1380
 30 ctctcactg cattttccat acaaagtgtt ctattttatt gttccttctt gtaataaagg 1440
 gaagataaaa ccaaaaaaaaa aaaaaaaaaa acggaattcc 1480

<210> 10

30 <211> 370

<212> PRT

<213> Homo sapiens

<220>

35 <223> Description of Sequence: N/A

<400> 10

Met Leu Gly Ala Val Glu Gly Pro Arg Trp Lys Gln Ala Glu Asp Ile

	1		5		10		15									
	Arg	Asp	Ile	Tyr	Asp	Phe	Arg	Asp	Val	Leu	Gly	Thr	Gly	Ala	Phe	Ser
			20				25						30			
	Glu	Val	Ile	Leu	Ala	Glu	Asp	Lys	Arg	Thr	Gln	Lys	Leu	Val	Ala	Ile
5			35				40					45				
	Lys	Cys	Ile	Ala	Lys	Glu	Ala	Leu	Glu	Gly	Lys	Glu	Gly	Ser	Met	Glu
		50				55					60					
	Asn	Glu	Ile	Ala	Val	Leu	His	Lys	Ile	Lys	His	Pro	Asn	Ile	Val	Ala
	65				70				75					80		
10	Leu	Asp	Asp	Ile	Tyr	Glu	Ser	Gly	Gly	His	Leu	Tyr	Leu	Ile	Met	Gln
			85					90					95			
	Leu	Val	Ser	Gly	Gly	Glu	Leu	Phe	Asp	Arg	Ile	Val	Glu	Lys	Gly	Phe
		100					105				110					
	Tyr	Thr	Glu	Arg	Asp	Ala	Ser	Arg	Leu	Ile	Phe	Gln	Val	Leu	Asp	Ala
15		115					120				125					
	Val	Lys	Tyr	Leu	His	Asp	Leu	Gly	Ile	Val	His	Arg	Asp	Leu	Lys	Pro
		130				135				140						
	Glu	Asn	Leu	Leu	Tyr	Tyr	Ser	Leu	Asp	Glu	Asp	Ser	Lys	Ile	Met	Ile
	145			150				155					160			
20	Ser	Asp	Phe	Gly	Leu	Ser	Lys	Met	Glu	Asp	Pro	Gly	Ser	Val	Leu	Ser
			165					170					175			
	Thr	Ala	Cys	Gly	Thr	Pro	Gly	Tyr	Val	Ala	Pro	Glu	Val	Leu	Ala	Gln
		180					185				190					
	Lys	Pro	Tyr	Ser	Lys	Ala	Val	Asp	Cys	Trp	Ser	Ile	Gly	Val	Ile	Ala
25		195					200				205					
	Tyr	Ile	Leu	Leu	Cys	Gly	Tyr	Pro	Pro	Phe	Tyr	Asp	Glu	Asn	Asp	Ala
		210				215				220						
	Lys	Leu	Phe	Glu	Gln	Ile	Leu	Lys	Ala	Glu	Tyr	Glu	Phe	Asp	Ser	Pro
	225			230				235					240			
30	Tyr	Trp	Asp	Asp	Ile	Ser	Asp	Ser	Ala	Lys	Asp	Phe	Ile	Arg	His	Leu
			245					250					255			
	Met	Glu	Lys	Asp	Pro	Glu	Lys	Arg	Phe	Thr	Cys	Glu	Gln	Ala	Leu	Gln
		260					265				270					
	His	Pro	Trp	Ile	Ala	Gly	Asp	Thr	Ala	Leu	Asp	Lys	Asn	Ile	His	Gln
35		275					280				285					
	Ser	Val	Ser	Glu	Gln	Ile	Lys	Lys	Asn	Phe	Ala	Lys	Ser	Lys	Trp	Lys
		290				295				300						
	Gln	Ala	Phe	Asn	Ala	Thr	Ala	Val	Val	Arg	His	Met	Arg	Lys	Leu	Gln

cactgttttg tatgaccccg ccgaagcaga agccccacca cctcaaattt atgatgccca 1080
 gttggaagaa agagaacatg caattgaaga atggaaagag ctaatttaca aagaagtcac 1140
 ggattgggaa gaaagaagca agaatggtgt tgtaaaagat cagccttcag atgcagcagt 1200
 aagtagcaac gccactcctt ctcatgtctt atcgatcaat gacatttcat ccatgtccac 1260
 5 tgagcagacg ctggcctcag acacagacag cagtcttgat gcctcgacgg gacccttga 1320
 aggctgtcga tgatagggtta gaaatagcaa acctgtcagc attgaaggaa ctctcacctc 1380
 cgtgggcctg aaatgcttgg gagttgatgg aaccaaatag aaaaactcca tgttctgcat 1440
 gtaagaaaca caatgccttg ccctattcag acctgatagg attgcctgct tagatgataa 1500
 aatgaggcag aatatgtctg aagaaaaaaaa ttgcaagcca cacttctaga gattttgttc 1560
 10 aagatcattt caggtgagca gttagagtag gtgaatttgt ttcaaattgt actagtgaca 1620
 gtttctcatc atctgtaact gttgagatgt atgtgcatgt gaccacaaat gcttgcttgg 1680
 acttgcccat ctagcacttt ggaaatcagt atttaaatgc caaataatct tccaggtagt 1740
 gctgcttctg aagttatctc ttaatcctct taagtaattt gg 1782

15

<210> 12

<211> 424

<212> PRT

<213> Homo sapiens

20

<220>

<223> Description of Sequence: N/A

<400> 12

25 Met Ser Asp Ser Lys Cys Asp Ser Gln Phe Tyr Ser Val Gln Val Ala
 1 5 10 15
 Asp Ser Thr Phe Thr Val Leu Lys Arg Tyr Gln Gln Leu Lys Pro Ile
 20 25 30
 Gly Ser Gly Ala Gln Gly Ile Val Cys Ala Ala Phe Asp Thr Val Leu
 30 35 40 45
 Gly Ile Ser Val Ala Val Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln
 50 55 60
 Thr His Ala Lys Arg Ala Tyr Arg Glu Leu Val Leu Leu Lys Cys Val
 65 70 75 80
 35 Asn His Lys Asn Ile Ile Ser Leu Leu Asn Val Phe Thr Pro Gln Lys
 85 90 95
 Thr Leu Glu Glu Phe Gln Asp Val Tyr Leu Val Met Glu Leu Met Asp
 100 105 110

Ala Asn Leu Cys Gln Val Ile His Met Glu Leu Asp His Glu Arg Met
 115 120 125
 Ser Tyr Leu Leu Tyr Gln Met Leu Cys Gly Ile Lys His Leu His Ser
 130 135 140
 5 Ala Gly Ile Ile His Arg Asp Leu Lys Pro Ser Asn Ile Val Val Lys
 145 150 155 160
 Ser Asp Cys Thr Leu Lys Ile Leu Asp Phe Gly Leu Ala Arg Thr Ala
 165 170 175
 Cys Thr Asn Phe Met Met Thr Pro Tyr Val Val Thr Arg Tyr Tyr Arg
 10 180 185 190
 Ala Pro Glu Val Ile Leu Gly Met Gly Tyr Lys Glu Asn Val Asp Ile
 195 200 205
 Trp Ser Val Gly Cys Ile Met Gly Glu Leu Val Lys Gly Cys Val Ile
 210 215 220
 15 Phe Gln Gly Thr Asp His Ile Asp Gln Trp Asn Lys Val Ile Glu Gln
 225 230 235 240
 Leu Gly Thr Pro Ser Ala Glu Phe Met Lys Lys Leu Gln Pro Thr Val
 245 250 255
 Arg Asn Tyr Val Glu Asn Arg Pro Lys Tyr Pro Gly Ile Lys Phe Glu
 20 260 265 270
 Glu Leu Phe Pro Asp Trp Ile Phe Pro Ser Glu Ser Glu Arg Asp Lys
 275 280 285
 Ile Lys Thr Ser Gln Ala Arg Asp Leu Leu Ser Lys Met Leu Val Ile
 290 295 300
 25 Asp Pro Asp Lys Arg Ile Ser Val Asp Glu Ala Leu Arg His Pro Tyr
 305 310 315 320
 Ile Thr Val Trp Tyr Asp Pro Ala Glu Ala Glu Ala Pro Pro Pro Gln
 325 330 335
 Ile Tyr Asp Ala Gln Leu Glu Glu Arg Glu His Ala Ile Glu Glu Trp
 30 340 345 350
 Lys Glu Leu Ile Tyr Lys Glu Val Met Asp Trp Glu Glu Arg Ser Lys
 355 360 365
 Asn Gly Val Val Lys Asp Gln Pro Ser Asp Ala Ala Val Ser Ser Asn
 370 375 380
 35 Ala Thr Pro Ser Gln Ser Ser Ser Ile Asn Asp Ile Ser Ser Met Ser
 385 390 395 400
 Thr Glu Gln Thr Leu Ala Ser Asp Thr Asp Ser Ser Leu Asp Ala Ser
 405 410 415

Thr Gly Pro Leu Glu Gly Cys Arg

420

5

<210> 13

<211> 3668

<212> DNA

<213> Homo sapiens

10

<220> LIMK-2

<223> Description of Sequence: N/A

<400> 13

15 gtggtcttcc cgcgctgag gcggcggcgg caggagctga ggggagttgt aggggaactga 60
ggggagctgc tgtgtcccc gcctcctcct cccatttcc gggctcccg gaccatgtcc 120
gcgctggcgg gtgaagatgt ctggaggtgt ccaggctgtg gggaccacat tgctccaagc 180
cagatatggt acaggactgt caacgaaacc tggcacggct cttgcttccg gtgttcagaa 240
tgccaggatt cctcaccaa ctggtactat gagaaggatg ggaagctcta ctgccccaaag 300
20 gactactggg ggaagtttg ggagttctgt catgggtgct cctgctgat gacagggcct 360
tttatggtgg ctggggagtt caagtaccac ccagagtgtc ttgcctgtat gagctgcaag 420
gtgatcattg aggatgggga tgcatatgca ctggtgcagc atgccaccct ctactgtggg 480
aagtgccaca atgaggtggt gctggcacc atgtttgaga gactctccac agagtctgtt 540
caggagcagc tgccctactc tgtcacgtc atctccatgc cggccaccac tgaaggcagg 600
25 cggggcttct cegtgtccgt ggagagtgcc tgctccaact acgccaccac tgtgcaagtg 660
aaagaggtca accggatgca catcagtccc aacaatcgaa acgccatcca ccctggggac 720
cgcatcctgg agatcaatgg gacccccgtc cgcacacttc gaggaggagg ggtggaggat 780
gcaattagcc agacgagcca gacacttcag ctgttgattg aacatgaccc cgtctcccaa 840
cgcttgacc agctgcggct ggaggcccgg ctgctcctc acatgcagaa tgccggacac 900
30 cccacgccc tcagcaccct ggacaccaag gagaatctgg aggggacact gaggagacgt 960
tccctaaggc gcagtaacag tatctccaag tccctggcc ccagctcccc aaaggagccc 1020
ctgctgttca gccgtgacat cagccgtca gaatccctc gttgttccag cagctattca 1080
cagcagatct tccggccctg tgacctaatc catggggagg tcctggggaa gggcttcttt 1140
gggcaggcta tcaagtgac acacaaagcc acgggcaaag tgatggtcat gaaagagtta 1200
35 attcgatgtg atgaggagac ccagaaaact tttctgactg aggtgaaagt gatgcgcagc 1260
ctggaccacc ccaatgtgct caagttcatt ggtgtgctgt acaaggataa gaagctgaac 1320
ctgctgacag agtacattga ggggggcaca ctgaaggact ttctgcgcag tatggatccg 1380
ttcccctggc agcagaaggt caggtttgcc aaaggaatcg cctccggaat ggcctatattg 1440

cactctatgt gcatcatcca ccgggatctg aactcgcaca actgcctcat caagttggac 1500
aagactgtgg tgggtggcaga ctttgggctg tcacggctca tagtggaaga gagggaaaagg 1560
gcccccatgg agaaggccac caccaagaaa cgcaccttgc gcaagaacga ccgcaagaag 1620
cgctacacgg tgggtgggaaa ccctactgg atggcccctg agatgctgaa cggaaagagc 1680
5 tatgatgaga cgggtggatat cttctccttt gggatcgttc tctgtgagat cattggggcag 1740
gtgtatgcag atcctgactg ccttccccga acactggact ttggcctcaa cgtgaagctt 1800
ttctgggaga agtttgttcc cacagattgt cccccggcct tcttcccgtt ggccgccatc 1860
tgctgcagac tggagcctga gagcagacca gcattctcga aattggagga ctcctttgag 1920
gccctctccc tgtacctggg ggagctgggc atcccgtgc ctgcagagct ggaggagtgtg 1980
10 gaccacactg tgagcatgca gtacggcctg acccgggact cacctcccta gccctggccc 2040
agccccctgc aggggggtgt tctacagcca gcattgcccc tctgtgcccc attcctgctg 2100
tgagcagggc cgtccgggct tctgtggat tggcggaatg tttagaagca gaacaagcca 2160
ttcctattac ctccccagga ggcaagtggg cgcagcacca gggaaatgta tctccacagg 2220
ttctggggcc tagttactgt ctgtaaatcc aatacttgcc tgaaagctgt gaagaagaaa 2280
15 aaaaccctg gcctttgggc caggaggaat ctgttactcg aatccacca ggaactccct 2340
ggcagtggat tgtgggaggc tcttgcttac actaatcagc gtgacctgga cctgctgggc 2400
aggatcccag ggtgaacctg cctgtgaact ctgaagtcac tagtccagct ggggtgcagga 2460
ggacttcaag tgtgtggacg aaagaaagac tgatggctca aagggtgtga aaaagtcagt 2520
gatgctcccc ctttctactc cagatcctgt ccttcttgga gcaagggtga gggagttagt 2580
20 tttgaagagt cccttaatat gtggtggaac aggccaggag ttagagaaag ggctggcttc 2640
tgtttacctg ctactggct ctagccagcc cagggaccac atcaatgtga gaggaagcct 2700
ccacctcatg ttttcaaact taatactgga gactggctga gaacttacgg acaacatcct 2760
ttctgtctga aacaaacagt cacaagcaca ggaagaggct gggggactag aaagaggccc 2820
tgccctctag aaagctcaga tcttggttcc tgttactcat actcgggtgg gctccttagt 2880
25 cagatgccta aaacattttg cctaaagctc gatgggttct ggaggacagt gtggcttgtc 2940
acaggcctag agtctgaggg aggggagtgg gagtctcagc aatctcttgg tcttggttcc 3000
atggcaacca ctgctcacc ttcaacatgc ctggtttagg cagcagcttg ggctgggaag 3060
aggtgggtggc agagtctcaa agctgagatg ctgagagaga tagtccctg agctgggcca 3120
tctgacttct acctcccatg tttgctctcc caactcatta gctcctgggc agcatcctcc 3180
30 tgagccacat gtgcaggtac tggaaaacct ccatcttggc tcccagagct ctaggaactc 3240
ttcatcacia ctagatttgc ctcttctaag tgtctatgag cttgcacat atttaataaa 3300
ttgggaatgg gtttggggta ttaatgcaat gtgtgggtgg tgtattggag cagggggaat 3360
tgataaagga gagtgggtgc tgttaatat atcttatcta ttgggtggta tgtgaaatat 3420
tgtacataga cctgatgagt tgtgggacca gatgtcatct ctggtcagag tttacttgct 3480
35 atatagactg tacttatgtg tgaagtttgc aagcttgctt tagggctgag ccctggactc 3540
ccagcagcag cacagttcag cattgtgtgg ctggttgttt cctggctgtc ccagcaagt 3600
gtaggagtgg tgggcctgaa ctgggccatt gatcagacta aataaattaa gcagttaaca 3660
taactggc 3668

<210> 14

<211> 638

5 <212> PRT

<213> Homo sapiens

<220>

<223> Description of Sequence: N/A

10

<400> 14

```

Met Ser Ala Leu Ala Gly Glu Asp Val Trp Arg Cys Pro Gly Cys Gly
  1              5              10              15
Asp His Ile Ala Pro Ser Gln Ile Trp Tyr Arg Thr Val Asn Glu Thr
15              20              25              30
Trp His Gly Ser Cys Phe Arg Cys Ser Glu Cys Gln Asp Ser Leu Thr
      35              40              45
Asn Trp Tyr Tyr Glu Lys Asp Gly Lys Leu Tyr Cys Pro Lys Asp Tyr
      50              55              60
20 Trp Gly Lys Phe Gly Glu Phe Cys His Gly Cys Ser Leu Leu Met Thr
      65              70              75              80
Gly Pro Phe Met Val Ala Gly Glu Phe Lys Tyr His Pro Glu Cys Phe
      85              90              95
Ala Cys Met Ser Cys Lys Val Ile Ile Glu Asp Gly Asp Ala Tyr Ala
25              100             105             110
Leu Val Gln His Ala Thr Leu Tyr Cys Gly Lys Cys His Asn Glu Val
      115             120             125
Val Leu Ala Pro Met Phe Glu Arg Leu Ser Thr Glu Ser Val Gln Glu
      130             135             140
30 Gln Leu Pro Tyr Ser Val Thr Leu Ile Ser Met Pro Ala Thr Thr Glu
      145             150             155             160
Gly Arg Arg Gly Phe Ser Val Ser Val Glu Ser Ala Cys Ser Asn Tyr
      165             170             175
Ala Thr Thr Val Gln Val Lys Glu Val Asn Arg Met His Ile Ser Pro
35              180             185             190
Asn Asn Arg Asn Ala Ile His Pro Gly Asp Arg Ile Leu Glu Ile Asn
      195             200             205
Gly Thr Pro Val Arg Thr Leu Arg Val Glu Glu Val Glu Asp Ala Ile

```


	210	215	220
	Ser Gln Thr	Ser Gln Thr Leu Gln Leu Leu Ile	Glu His Asp Pro Val
	225	230	235 240
	Ser Gln Arg Leu Asp	Gln Leu Arg Leu Glu Ala Arg Leu Ala Pro His	
5	245	250	255
	Met Gln Asn Ala Gly His Pro His Ala Leu Ser Thr Leu Asp Thr Lys		
	260	265	270
	Glu Asn Leu Glu Gly Thr Leu Arg Arg Arg Ser Leu Arg Arg Ser Asn		
	275	280	285
10	Ser Ile Ser Lys Ser Pro Gly Pro Ser Ser Pro Lys Glu Pro Leu Leu		
	290	295	300
	Phe Ser Arg Asp Ile Ser Arg Ser Glu Ser Leu Arg Cys Ser Ser Ser		
	305	310	315 320
	Tyr Ser Gln Gln Ile Phe Arg Pro Cys Asp Leu Ile His Gly Glu Val		
15	325	330	335
	Leu Gly Lys Gly Phe Phe Gly Gln Ala Ile Lys Val Thr His Lys Ala		
	340	345	350
	Thr Gly Lys Val Met Val Met Lys Glu Leu Ile Arg Cys Asp Glu Glu		
	355	360	365
20	Thr Gln Lys Thr Phe Leu Thr Glu Val Lys Val Met Arg Ser Leu Asp		
	370	375	380
	His Pro Asn Val Leu Lys Phe Ile Gly Val Leu Tyr Lys Asp Lys Lys		
	385	390	395 400
	Leu Asn Leu Leu Thr Glu Tyr Ile Glu Gly Gly Thr Leu Lys Asp Phe		
25	405	410	415
	Leu Arg Ser Met Asp Pro Phe Pro Trp Gln Gln Lys Val Arg Phe Ala		
	420	425	430
	Lys Gly Ile Ala Ser Gly Met Ala Tyr Leu His Ser Met Cys Ile Ile		
	435	440	445
30	His Arg Asp Leu Asn Ser His Asn Cys Leu Ile Lys Leu Asp Lys Thr		
	450	455	460
	Val Val Val Ala Asp Phe Gly Leu Ser Arg Leu Ile Val Glu Glu Arg		
	465	470	475 480
	Lys Arg Ala Pro Met Glu Lys Ala Thr Thr Lys Lys Arg Thr Leu Arg		
35	485	490	495
	Lys Asn Asp Arg Lys Lys Arg Tyr Thr Val Val Gly Asn Pro Tyr Trp		
	500	505	510
	Met Ala Pro Glu Met Leu Asn Gly Lys Ser Tyr Asp Glu Thr Val Asp		

	515	520	525
	Ile Phe Ser Phe Gly Ile Val Leu Cys Glu Ile Ile Gly Gln Val Tyr		
	530	535	540
	Ala Asp Pro Asp Cys Leu Pro Arg Thr Leu Asp Phe Gly Leu Asn Val		
5	545	550	555
	Lys Leu Phe Trp Glu Lys Phe Val Pro Thr Asp Cys Pro Pro Ala Phe		
	565	570	575
	Phe Pro Leu Ala Ala Ile Cys Cys Arg Leu Glu Pro Glu Ser Arg Pro		
	580	585	590
10	Ala Phe Ser Lys Leu Glu Asp Ser Phe Glu Ala Leu Ser Leu Tyr Leu		
	595	600	605
	Gly Glu Leu Gly Ile Pro Leu Pro Ala Glu Leu Glu Glu Leu Asp His		
	610	615	620
	Thr Val Ser Met Gln Tyr Gly Leu Thr Arg Asp Ser Pro Pro		
15	625	630	635

20 <210> 15
 <211> 2169
 <212> DNA
 <213> Homo sapiens

25 <220> PRK
 <223> Description of Sequence: N/A

30 <400> 15
 ccgcctccga gtgccttgcg cggacctgag ctggagatgc tggccgggct accgacgtca 60
 gaccccgggc gcctcatcac ggacccgcgc agcggccgca cctacctcaa aggccgcttg 120
 ttgggcaagg ggggcttcgc ccgctgctac gaggccactg acacagagac tggcagcgcc 180
 tacgctgtca aagtcatccc gcagagccgc gtcgccaagc cgcatacagc cgagaagatc 240
 ctaaattgaga ttgagctgca ccgagacctg cagcaccgcc acatcgtgcg tttttcgcac 300
 cactttgagg acgctgacaa catctacatt ttcttgagac tctgcagccg aaagtccctg 360
 gccacatctt ggaaggcccg gcacaccctg ttggagccag aagtgcgcta ctacctgcgg 420
 35 cagatccttt ctggcctcaa gtacttgac cagcgcgga tcttgaccg ggacctcaag 480
 ttgggaaatt ttttcatcac tgagaacatg gaactgaagg tgggggattt tgggctggca 540
 gcccggttgg agcctccgga gcagaggaag aagaccatct gtggcacccc caactatgtg 600
 gctccagaag tgctgctgag acagggccac ggccctgaag cggatgtatg gtcactgggc 660

5 tgtgtcatgt acacgctgct ctgcgggagc cctccctttg agacggctga cctgaaggag 720
 acgtaccgct gcatcaagca ggttcaactac acgctgcctg ccagcctctc actgcctgcc 780
 cggcagctcc tggccgcat ccttcgggccc tcaccccgag accgcccctc tattgaccag 840
 atcctgcgcc atgacttctt taccaagggc tacacccccg atcgactccc tatcagcagc 900
 10 tgcgtgacag tcccagacct gacaccccc aaccagcta ggagtctgtt tgccaaagtt 960
 accaagagcc tctttggcag aaagaagaag agtaagaatc atgcccagga gagggatgag 1020
 gtctccggtt tggtagcggt cctcatgcgc acatccgttg gccatcagga tgccaggcca 1080
 gaggtccag cagcttcttg cccagcccct gtcagcctgg tagagacagc acctgaagac 1140
 agctcacccc gtgggacact ggcaagcagt ggagatggat ttgaagaagg tctgactgtg 1200
 15 gccacagtag tggagtcagc cctttgtgct ctgagaaatt gtatagcttt catgccccca 1260
 gcggaacaga acccggtccc cctggcccag ccagagcctc tgggtgtgggt cagcaagtgg 1320
 gttgactact ccaataagtt cggttttggg tatcaactgt ccagccgccc tgtggctgtg 1380
 ctcttcaacg atggcacaca tatggccctg tcggccaaca gaaagactgt gcactacaat 1440
 cccaccagca caaagcactt ctccttctcc gtgggtgctg tgccccgggc cctgcagcct 1500
 20 cagctgggta tcctgcggta cttegcctcc tacatggagc agcacctcat gaaggggtgga 1560
 gatctgcccc gtgtggaaga ggtagaggta cctgctccgc ccttgcctgt gcagtgggtc 1620
 aagacggatc aggtctctct catgctgttt agtgatggca ctgtccaggt gaacttctac 1680
 ggggaccaca ccaagctgat tctcagtggc tgggagcccc tccttgtgac ttttgtggcc 1740
 cgaaatcgta gtgcttgtac ttacctcgct tcccaccttc ggcagctggg ctgctctcca 1800
 25 gacctgcggc agcgactccg ctatgctctg cgctgctcc gggaccgcag cccagcttag 1860
 gacccaagcc ctgaaggcct gaggcctgtg cctgtcaggg tctggccctt gcctttgtgg 1920
 ccttccccct tcctttgggt cctcactggg ggctttgggc cgaatcccc agggaaatcag 1980
 ggaccagctt tactggagtt gggggcggct tgtcttctgt ggctcctacc ccatctccaa 2040
 gataagcctg agccttagct cccagctagg gggcggtatt tatggaccac ttttatttat 2100
 30 tgtcagacac ttattttatt ggatgtgagc cccagggggc ctctcctag gataataaac 2160
 aattttgca 2169

<210> 16

30 <211> 607

<212> PRT

<213> Homo sapiens

<220>

35 <223> Description of Sequence: N/A

<400> 16

Met Leu Ala Gly Leu Pro Thr Ser Asp Pro Gly Arg Leu Ile Thr Asp

	1		5		10		15									
	Pro	Arg	Ser	Gly	Arg	Thr	Tyr	Leu	Lys	Gly	Arg	Leu	Leu	Gly	Lys	Gly
				20				25						30		
	Gly	Phe	Ala	Arg	Cys	Tyr	Glu	Ala	Thr	Asp	Thr	Glu	Thr	Gly	Ser	Ala
5			35					40						45		
	Tyr	Ala	Val	Lys	Val	Ile	Pro	Gln	Ser	Arg	Val	Ala	Lys	Pro	His	Gln
			50					55						60		
	Arg	Glu	Lys	Ile	Leu	Asn	Glu	Ile	Glu	Leu	His	Arg	Asp	Leu	Gln	His
			65					70						75		
10	Arg	His	Ile	Val	Arg	Phe	Ser	His	His	Phe	Glu	Asp	Ala	Asp	Asn	Ile
					85					90					95	
	Tyr	Ile	Phe	Leu	Glu	Leu	Cys	Ser	Arg	Lys	Ser	Leu	Ala	His	Ile	Trp
				100						105					110	
	Lys	Ala	Arg	His	Thr	Leu	Leu	Glu	Pro	Glu	Val	Arg	Tyr	Tyr	Leu	Arg
15				115						120					125	
	Gln	Ile	Leu	Ser	Gly	Leu	Lys	Tyr	Leu	His	Gln	Arg	Gly	Ile	Leu	His
				130						135					140	
	Arg	Asp	Leu	Lys	Leu	Gly	Asn	Phe	Phe	Ile	Thr	Glu	Asn	Met	Glu	Leu
				145						150					155	
20	Lys	Val	Gly	Asp	Phe	Gly	Leu	Ala	Ala	Arg	Leu	Glu	Pro	Pro	Glu	Gln
					165					170					175	
	Arg	Lys	Lys	Thr	Ile	Cys	Gly	Thr	Pro	Asn	Tyr	Val	Ala	Pro	Glu	Val
					180					185					190	
	Leu	Leu	Arg	Gln	Gly	His	Gly	Pro	Glu	Ala	Asp	Val	Trp	Ser	Leu	Gly
25					195					200					205	
	Cys	Val	Met	Tyr	Thr	Leu	Leu	Cys	Gly	Ser	Pro	Pro	Phe	Glu	Thr	Ala
					210					215					220	
	Asp	Leu	Lys	Glu	Thr	Tyr	Arg	Cys	Ile	Lys	Gln	Val	His	Tyr	Thr	Leu
					225					230					235	
30	Pro	Ala	Ser	Leu	Ser	Leu	Pro	Ala	Arg	Gln	Leu	Leu	Ala	Ala	Ile	Leu
					245					250					255	
	Arg	Ala	Ser	Pro	Arg	Asp	Arg	Pro	Ser	Ile	Asp	Gln	Ile	Leu	Arg	His
					260					265					270	
	Asp	Phe	Phe	Thr	Lys	Gly	Tyr	Thr	Pro	Asp	Arg	Leu	Pro	Ile	Ser	Ser
35					275					280					285	
	Cys	Val	Thr	Val	Pro	Asp	Leu	Thr	Pro	Pro	Asn	Pro	Ala	Arg	Ser	Leu
					290					295					300	
	Phe	Ala	Lys	Val	Thr	Lys	Ser	Leu	Phe	Gly	Arg	Lys	Lys	Lys	Ser	Lys

	305		310		315		320									
	Asn	His	Ala	Gln	Glu	Arg	Asp	Glu	Val	Ser	Gly	Leu	Val	Ser	Gly	Leu
				325						330					335	
	Met	Arg	Thr	Ser	Val	Gly	His	Gln	Asp	Ala	Arg	Pro	Glu	Ala	Pro	Ala
5				340						345					350	
	Ala	Ser	Gly	Pro	Ala	Pro	Val	Ser	Leu	Val	Glu	Thr	Ala	Pro	Glu	Asp
				355						360					365	
	Ser	Ser	Pro	Arg	Gly	Thr	Leu	Ala	Ser	Ser	Gly	Asp	Gly	Phe	Glu	Glu
				370						375					380	
10	Gly	Leu	Thr	Val	Ala	Thr	Val	Val	Glu	Ser	Ala	Leu	Cys	Ala	Leu	Arg
				385						390					395	
	Asn	Cys	Ile	Ala	Phe	Met	Pro	Pro	Ala	Glu	Gln	Asn	Pro	Ala	Pro	Leu
					405						410				415	
	Ala	Gln	Pro	Glu	Pro	Leu	Val	Trp	Val	Ser	Lys	Trp	Val	Asp	Tyr	Ser
15				420						425					430	
	Asn	Lys	Phe	Gly	Phe	Gly	Tyr	Gln	Leu	Ser	Ser	Arg	Arg	Val	Ala	Val
				435						440					445	
	Leu	Phe	Asn	Asp	Gly	Thr	His	Met	Ala	Leu	Ser	Ala	Asn	Arg	Lys	Thr
				450						455					460	
20	Val	His	Tyr	Asn	Pro	Thr	Ser	Thr	Lys	His	Phe	Ser	Phe	Ser	Val	Gly
				465						470					475	
	Ala	Val	Pro	Arg	Ala	Leu	Gln	Pro	Gln	Leu	Gly	Ile	Leu	Arg	Tyr	Phe
					485					490					495	
	Ala	Ser	Tyr	Met	Glu	Gln	His	Leu	Met	Lys	Gly	Gly	Asp	Leu	Pro	Ser
25				500						505					510	
	Val	Glu	Glu	Val	Glu	Val	Pro	Ala	Pro	Pro	Leu	Leu	Leu	Gln	Trp	Val
				515						520					525	
	Lys	Thr	Asp	Gln	Ala	Leu	Leu	Met	Leu	Phe	Ser	Asp	Gly	Thr	Val	Gln
				530						535					540	
30	Val	Asn	Phe	Tyr	Gly	Asp	His	Thr	Lys	Leu	Ile	Leu	Ser	Gly	Trp	Glu
				545						550					555	
	Pro	Leu	Leu	Val	Thr	Phe	Val	Ala	Arg	Asn	Arg	Ser	Ala	Cys	Thr	Tyr
					565					570					575	
	Leu	Ala	Ser	His	Leu	Arg	Gln	Leu	Gly	Cys	Ser	Pro	Asp	Leu	Arg	Gln
35				580						585					590	
	Arg	Leu	Arg	Tyr	Ala	Leu	Arg	Leu	Leu	Arg	Asp	Arg	Ser	Pro	Ala	
				595						600					605	

<210> 17

<211> 3492

5 <212> DNA

<213> Homo sapiens

<220> PTP-SL

<223> Description of Sequence: N/A

10

<400> 17

15

20

25

30

35

cagctaagac ccggagaggt ggaatttcac ttgaaattc ccttgccctcg tgaggggccgg 60
cgctgggcat gctcagtagc cgcggcgctg ctgctgggct gctgggctgg cgcggagtc 120
accctgccgt ctccgccttg gcttctgggc gtccagaagg ccaggcattt gccgcctctg 180
agcgcttctg ttcccccttac ccgcaacctc ctactgctct tcctctctcc ctctcttagg 240
gaggttgaag ctggtgctgg tttctgtcgg cgccacagac tgactgctct gcaaacccca 300
gccgaggacc tgaatcccgg agactagaag acccttggcg gtggctcttt ctaatagcac 360
tttacctgaa gtgggggtcgt ggtggagttt ctctccacc tctcaatgca aacactatgc 420
ggagagcagt ctgcttccct gcgctgtgcc tgctccttaa tcttcacgct gcagggtgct 480
tttcaggaaa caatgatcat tttttggcaa ttaatcagaa gaagagtggg aagccggtat 540
tcatttataa gcattcacaa gacattgaga agagcctgga tatagcccca caaaaaatct 600
acagacatag ctaccattcc tcttccgaag ctcaagtaag caaacgccac cagattgtca 660
attcagcatt tcctagaccc gcatatgacc cgtctctcaa tctgctggcc atggatggtc 720
aagatcttga agtgaaaaat ctcccaatcc cagcagcaaa tgtaattgtg gtgacactgc 780
aaatggatgt aaacaagctg aacataacct tgcttcggat cttccgcca ggagtggctg 840
cagcttttagg actcttacct cagcaagtgc acatcaatcg cctcattgga aagaagaaca 900
gtattgaact gtttgtgtct ccataaacc gaaaaacagg aatttctgat gctctgccct 960
ctgaggaagt tcttcgttca cttaatatca atgttttgca tcaaagttta tcccagtttg 1020
gaattacaga agtctctcct gagaaaaatg ttttacaagg gcagcatgaa gcggacaaaa 1080
tctggagcaa agaaggattt tatgctgttg tcatttttct cagcatcttt gttattatag 1140
taacgtgttt gatgattctt tacagattaa aagaaagatt tcagctttcc ttaagacaag 1200
acaaagagaa aaaccaggag atccacctat cgcccatcac attacagcca gcactgtccg 1260
aggcaaagac agtccacagc atggtccaac ctgagcaggc cccaaaggta ctgaatgttg 1320
tcgtggaccc tcaaggccga ggtgctcctg agatcagagc taccaccgct acctctgttt 1380
gcccttctcc tttcaaaatg aagcccatag gacttcaaga gagaagaggg tccaacgtat 1440
ctcttacatt ggacatgagt agcttgggga acattgaacc ctttgtgtct ataccaacac 1500
cacgggagaa ggtagcaatg gagtatctgc agtcagccag ccgaattctc acaaggctctc 1560
agctgagggga cgtcgtggca agttcacatt tactccaaag tgaattcatg gaaataccga 1620

tgaactttgt ggatcccaaa gaaattgata ttccgcgtca tggaactaaa aatcgctata 1680
 agaccatttt accaaatccc ctacagcagag tgtgtttaag accaaaaaat gtaaccgatt 1740
 cattgagcac ctacattaat gctaattata ttaggggcta cagtggcaag gagaaagcct 1800
 tcattgccac gcagggcccc atgatcaaca ccgtggatga tttctggcag atggtttggc 1860
 5 aggaagacag ccctgtgatt gttatgatca caaaactcaa agaaaaaat gagaaatgtg 1920
 tgctatactg gccggaaaag agagggatat atggaaaagt tgaggttctg gttatcagtg 1980
 taaatgaatg tgataactac accattcgaa accttgtctt aaagcaagga agccacaccc 2040
 aacatgtgaa gcattactgg tacacctcat ggctgatca caagactcca gacagtgcc 2100
 agccccctct acagctcatg ctggatgtag aagaagacag acttgcttcc cagggccgag 2160
 10 ggctgtggt tgtccactgc agtgcaggaa taggtagaac aggggtgtttt attgctacat 2220
 ccattggctg tcaacagctg aaagaagaag gagttgtgga tgcactaagc attgtctgcc 2280
 agcttcgtat ggatagaggt ggaatggctc aaaccagtga gcagtatgaa tttgtgcacc 2340
 atgctctgtg cctgtatgag agcagacttt cagcagagac tgtccagtga gtcattgaag 2400
 acttgtcaga ccatcaatct cttgggggtga ttaatcaaat taccaccca aggcttctag 2460
 15 aaggagcttc ctgcaatgga aggaaggaga agctctgaag cccatgtatg gcatggattg 2520
 tggaagactg ggcaacatat ttaagatttc cagctccttg tgtatatgaa tgcatttcta 2580
 agcatcccc aaattattct gaaggttttt tgatgatgga ggtatgatag gttatcaca 2640
 cagcctaagg cagattttgt tttgtctgta ctgactctat ctgccacaca gaatgtatgt 2700
 atgtaatat cagtaataaa tgcatcagg tgatgactgg atgagctgct gaagacattc 2760
 20 gtattatgtg ttagatgctt taatgtttgc aaaatctgcc ttgtgaatgg actgtcagct 2820
 gttaaactgt tcctgttttg aagtgtatt acctttctca gttaccagaa tcttgctgct 2880
 aaagttgcaa gtgattgata atggattttt aacagagaag tctttgtttt tgaaaaacaa 2940
 aaatcaaaaa cagtaactat tttatatgga aatgtgtctt gataatatta cctattaaat 3000
 gtgtatttat agtccctcct atcaacaat tacagagcac aatgattgtc attgggtata 3060
 25 tatgtattta ctctctatta ttgggcataa aggtggcttc tgctccagaa ctctatccac 3120
 tgtatttcca catcgtgagt cattttactt taaaaggga aaacaaattt gtagcaactc 3180
 tgaagtatca agagttttaa ctacttgact ctcttttgct aagaagggat ttttgaatat 3240
 gctatctacc tggaatctct ctctcaacaa aaggatatat ccttcaggaa tgatataatc 3300
 tgtcccatth tcgaggctcc ttataaggac atttccatgt atgtccttac atttctgaaa 3360
 30 gctttcaatc ttcaagagcc aaaaaaatt aaaataacta ccctcagcaa aactagctg 3420
 ttctgctcat atatgaattt ttaatgcagc aatgttgact ttgtttcata ctgccaataa 3480
 actcttaata ct 3492

35 <210> 18
 <211> 657
 <212> PRT
 <213> Homo sapiens

<220>

<223> Description of Sequence: N/A

5 <400> 18

Met Arg Arg Ala Val Cys Phe Pro Ala Leu Cys Leu Leu Leu Asn Leu

1 5 10 15

His Ala Ala Gly Cys Phe Ser Gly Asn Asn Asp His Phe Leu Ala Ile

20 25 30

10 Asn Gln Lys Lys Ser Gly Lys Pro Val Phe Ile Tyr Lys His Ser Gln

35 40 45

Asp Ile Glu Lys Ser Leu Asp Ile Ala Pro Gln Lys Ile Tyr Arg His

50 55 60

Ser Tyr His Ser Ser Ser Glu Ala Gln Val Ser Lys Arg His Gln Ile

15 65 70 75 80

Val Asn Ser Ala Phe Pro Arg Pro Ala Tyr Asp Pro Ser Leu Asn Leu

85 90 95

Leu Ala Met Asp Gly Gln Asp Leu Glu Val Glu Asn Leu Pro Ile Pro

100 105 110

20 Ala Ala Asn Val Ile Val Val Thr Leu Gln Met Asp Val Asn Lys Leu

115 120 125

Asn Ile Thr Leu Leu Arg Ile Phe Arg Gln Gly Val Ala Ala Ala Leu

130 135 140

Gly Leu Leu Pro Gln Gln Val His Ile Asn Arg Leu Ile Gly Lys Lys

25 145 150 155 160

Asn Ser Ile Glu Leu Phe Val Ser Pro Ile Asn Arg Lys Thr Gly Ile

165 170 175

Ser Asp Ala Leu Pro Ser Glu Glu Val Leu Arg Ser Leu Asn Ile Asn

180 185 190

30 Val Leu His Gln Ser Leu Ser Gln Phe Gly Ile Thr Glu Val Ser Pro

195 200 205

Glu Lys Asn Val Leu Gln Gly Gln His Glu Ala Asp Lys Ile Trp Ser

210 215 220

Lys Glu Gly Phe Tyr Ala Val Val Ile Phe Leu Ser Ile Phe Val Ile

35 225 230 235 240

Ile Val Thr Cys Leu Met Ile Leu Tyr Arg Leu Lys Glu Arg Phe Gln

245 250 255

Leu Ser Leu Arg Gln Asp Lys Glu Lys Asn Gln Glu Ile His Leu Ser

	260	265	270
	Pro Ile Thr Leu Gln Pro Ala Leu Ser Glu Ala Lys Thr Val His Ser		
	275	280	285
	Met Val Gln Pro Glu Gln Ala Pro Lys Val Leu Asn Val Val Val Asp		
5	290	295	300
	Pro Gln Gly Arg Gly Ala Pro Glu Ile Arg Ala Thr Thr Ala Thr Ser		
	305	310	315
	Val Cys Pro Ser Pro Phe Lys Met Lys Pro Ile Gly Leu Gln Glu Arg		
	325	330	335
10	Arg Gly Ser Asn Val Ser Leu Thr Leu Asp Met Ser Ser Leu Gly Asn		
	340	345	350
	Ile Glu Pro Phe Val Ser Ile Pro Thr Pro Arg Glu Lys Val Ala Met		
	355	360	365
	Glu Tyr Leu Gln Ser Ala Ser Arg Ile Leu Thr Arg Ser Gln Leu Arg		
15	370	375	380
	Asp Val Val Ala Ser Ser His Leu Leu Gln Ser Glu Phe Met Glu Ile		
	385	390	395
	Pro Met Asn Phe Val Asp Pro Lys Glu Ile Asp Ile Pro Arg His Gly		
	405	410	415
20	Thr Lys Asn Arg Tyr Lys Thr Ile Leu Pro Asn Pro Leu Ser Arg Val		
	420	425	430
	Cys Leu Arg Pro Lys Asn Val Thr Asp Ser Leu Ser Thr Tyr Ile Asn		
	435	440	445
	Ala Asn Tyr Ile Arg Gly Tyr Ser Gly Lys Glu Lys Ala Phe Ile Ala		
25	450	455	460
	Thr Gln Gly Pro Met Ile Asn Thr Val Asp Asp Phe Trp Gln Met Val		
	465	470	475
	Trp Gln Glu Asp Ser Pro Val Ile Val Met Ile Thr Lys Leu Lys Glu		
	485	490	495
30	Lys Asn Glu Lys Cys Val Leu Tyr Trp Pro Glu Lys Arg Gly Ile Tyr		
	500	505	510
	Gly Lys Val Glu Val Leu Val Ile Ser Val Asn Glu Cys Asp Asn Tyr		
	515	520	525
	Thr Ile Arg Asn Leu Val Leu Lys Gln Gly Ser His Thr Gln His Val		
35	530	535	540
	Lys His Tyr Trp Tyr Thr Ser Trp Pro Asp His Lys Thr Pro Asp Ser		
	545	550	555
	Ala Gln Pro Leu Leu Gln Leu Met Leu Asp Val Glu Glu Asp Arg Leu		

565 570 575
 Ala Ser Gln Gly Arg Gly Pro Val Val Val His Cys Ser Ala Gly Ile
 580 585 590
 Gly Arg Thr Gly Cys Phe Ile Ala Thr Ser Ile Gly Cys Gln Gln Leu
 5 595 600 605
 Lys Glu Glu Gly Val Val Asp Ala Leu Ser Ile Val Cys Gln Leu Arg
 610 615 620
 Met Asp Arg Gly Gly Met Val Gln Thr Ser Glu Gln Tyr Glu Phe Val
 625 630 635 640
 10 His His Ala Leu Cys Leu Tyr Glu Ser Arg Leu Ser Ala Glu Thr Val
 645 650 655
 Gln

 15
 <210> 19
 <211> 985
 <212> DNA
 <213> Homo sapiens
 20
 <220> HSP86
 <223> Description of Sequence: N/A

 <400> 19
 25 ccggcccggt gtggctgtgc cgttggctcct gtgcggtcac ttagccaaga tgcctgagga 60
 aaccagacc caagaccaac cgatggagga ggaggaggtt gagacgttcg cctttcaggc 120
 agaaattgcc cagttgatgt cattgatcat caatactttc tactcgaaca aagagatctt 180
 tctgagagag ctcatattca attcatcaga tgcattggac aaaatccggt atgaaagctt 240
 gacagatccc agtaaattag actctgggaa agagctgcat attaacctta taccgaacaa 300
 30 acaagatcga actctcacta ttgtggatac tggaattgga atgaccaagg ctgacttgat 360
 caataacctt ggtactatcg ccaagtctgg gaccaaagcg ttcattggaag ctttgcaggc 420
 tgggtgcagat atctctatga ttggccagtt cgggtgttgg ttttattctg cttatttggt 480
 tgctgagaaa gtaactgtga tcaccaaaca taacgatgat gagcagtacg cttgggagtc 540
 ctgagcaggg ggatcattca cagtggaggac agacacaggt gaacctatgg gtcgtggaac 600
 35 aaaagttatc ctacacctga aagaagacca aactgagtac ttggaggaac gaagaataaa 660
 ggagattgtg aagaaacatt ctgagtttat tggatatccc attactcttt ttgtggagaa 720
 ggaacgtgat aaagaagtaa gcgatgatga ggctgaagaa aaggaagaca aagaagaaga 780
 aaaagaaaaa gaagagaaag agtcggaaga caaacctgaa attgaagatg ttggttctga 840

tgaggaagaa gaaaagaagg atggtgacaa gaagaagaag aagaagatta aggaaaagta 900
catcgatcaa gaagagctca acaaaacaaa gcccatctgg accagaaatc ccgacgatat 960
tactaatgag gagtacggag aattc 985

5

<210> 20
<211> 312
<212> PRT
<213> Homo sapiens

10

<220>
<223> Description of Sequence: N/A

<400> 20

15 Met Pro Glu Glu Thr Gln Thr Gln Asp Gln Pro Met Glu Glu Glu Glu
1 5 10 15

Val Glu Thr Phe Ala Phe Gln Ala Glu Ile Ala Gln Leu Met Ser Leu
20 25 30

20

Ile Ile Asn Thr Phe Tyr Ser Asn Lys Glu Ile Phe Leu Arg Glu Leu
35 40 45

25

Ile Ser Asn Ser Ser Asp Ala Leu Asp Lys Ile Arg Tyr Glu Ser Leu
50 55 60

Thr Asp Pro Ser Lys Leu Asp Ser Gly Lys Glu Leu His Ile Asn Leu
65 70 75 80

30

Ile Pro Asn Lys Gln Asp Arg Thr Leu Thr Ile Val Asp Thr Gly Ile
85 90 95

Gly Met Thr Lys Ala Asp Leu Ile Asn Asn Leu Gly Thr Ile Ala Lys
100 105 110

35

Ser Gly Thr Lys Ala Phe Met Glu Ala Leu Gln Ala Gly Ala Asp Ile
115 120 125

Ser Met Ile Gly Gln Phe Gly Val Gly Phe Tyr Ser Ala Tyr Leu Val
130 135 140

Ala Glu Lys Val Thr Val Ile Thr Lys His Asn Asp Asp Glu Gln Tyr
5 145 150 155 160

Ala Trp Glu Ser Ser Ala Gly Gly Ser Phe Thr Val Arg Thr Asp Thr
165 170 175

10 Gly Glu Pro Met Gly Arg Gly Thr Lys Val Ile Leu His Leu Lys Glu
180 185 190

Asp Gln Thr Glu Tyr Leu Glu Glu Arg Arg Ile Lys Glu Ile Val Lys
195 200 205

15 Lys His Ser Gln Phe Ile Gly Tyr Pro Ile Thr Leu Phe Val Glu Lys
210 215 220

Glu Arg Asp Lys Glu Val Ser Asp Asp Glu Ala Glu Glu Lys Glu Asp
20 225 230 235 240

Lys Glu Glu Glu Lys Glu Lys Glu Glu Lys Glu Ser Glu Asp Lys Pro
245 250 255

25 Glu Ile Glu Asp Val Gly Ser Asp Glu Glu Glu Glu Lys Lys Asp Gly
260 265 270

Asp Lys Lys Lys Lys Lys Lys Ile Lys Glu Lys Tyr Ile Asp Gln Glu
275 280 285

30 Glu Leu Asn Lys Thr Lys Pro Ile Trp Thr Arg Asn Pro Asp Asp Ile
290 295 300

Thr Asn Glu Glu Tyr Gly Glu Phe
35 305 310

(19) World Intellectual Property Organization
International Bureau(43) International Publication Date
21 November 2002 (21.11.2002)

PCT

(10) International Publication Number
WO 02/093164 A3(51) International Patent Classification⁷: **A61K 31/506**,
G01N 33/68, C12Q 1/42, 1/48, C07K 16/40, C12N 15/11,
A61P 25/28, C07D 401/04, 409/14, 401/14, 471/04(74) Agents: **LEIDESCHER, Thomas et al.**; Zimmermann &
Partner, Postfach 330 920, 80069 München (DE).

(21) International Application Number: PCT/EP02/05420

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU,
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,
MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG,
SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ,
VN, YU, ZA, ZM, ZW.

(22) International Filing Date: 16 May 2002 (16.05.2002)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
01111858.5 16 May 2001 (16.05.2001) EP
60/293,528 29 May 2001 (29.05.2001) US
01117113.9 13 July 2001 (13.07.2001) EP
60/305,898 18 July 2001 (18.07.2001) US(84) Designated States (*regional*): ARIPO patent (GH, GM,
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),
Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),
European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR,
GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent
(BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR,
NE, SN, TD, TG).(71) Applicant (*for all designated States except US*): **AXXIMA
PHARMACEUTICALS AG** [DE/DE]; Am Klopferspitz
19, 82152 Martinsried (DE).

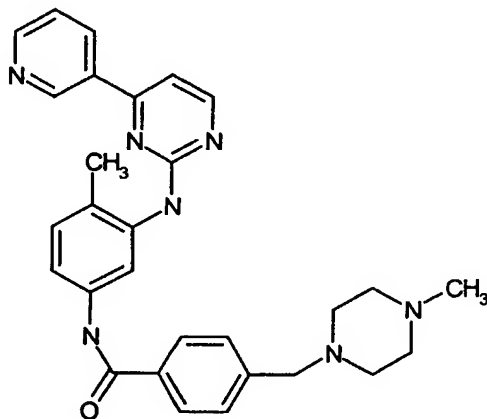
Published:

— with international search report

(72) Inventors; and

(75) Inventors/Applicants (*for US only*): **STEIN-GER-
LACH, Matthias** [DE/DE]; Stockdorfer Strasse 38A,
81475 München (DE). **SALASSIDIS, Konstadinos**
[GR/DE]; Echinger strasse 20, 85386 Eching (DE).
BACHER, Gerald [DE/DE]; Kriegerstrasse 62, 82110
Germering (DE). **MÜLLER, Stefan** [DE/DE]; Thalkirch-
ner Str. 184, 81371 München (DE).(88) Date of publication of the international search report:
4 September 2003*For two-letter codes and other abbreviations, refer to the "Guid-
ance Notes on Codes and Abbreviations" appearing at the begin-
ning of each regular issue of the PCT Gazette.*

(54) Title: PYRIDYLPYRIMIDINE DERIVATIVES AS EFFECTIVE COMPOUNDS AGAINST PRION DISEASES

Compound 53 (GleevecTM)(57) Abstract: The present invention relates to pyridylpyrim-
idine derivatives of the general formula (I) : wherein R
represents hydrogen or methyl and Z represents nitrogen
containing functional groups, the use of the pyridylpyrimidine
derivatives as pharmaceutically active agents, especially
for the prophylaxis and/or treatment of prion infections and
prion diseases, as well as compositions containing at least
one pyridylpyrimidine derivative and/or pharmaceutically
acceptable salt thereof. Furthermore, the present invention
is directed to methods for preventing and/or treating prion
infections and prion diseases using said pyridylpyrimidine
derivatives. Human cellular protein kinases, phosphatases and
cellular signal transduction molecules are disclosed as targets
for detecting, preventing and/or treating prion infections
and diseases, especially BSE, vCJD, or CJD, which can be
inhibited by the inventive pyridylpyrimidine derivatives.

WO 02/093164 A3

INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 02/05420

A. CLASSIFICATION OF SUBJECT MATTER IPC 7 A61K31/506 G01N33/68 C12Q1/42 C12Q1/48 C07K16/40 C12N15/11 A61P25/28 C07D401/04 C07D409/14 C07D401/14 C07D471/04		
According to International Patent Classification (IPC) or to both national classification and IPC		
B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) IPC 7 C12N C07D		
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched		
Electronic data base consulted during the international search (name of data base and, where practical, search terms used)		
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	ZIMMERMANN J ET AL: "Phenylamino-pyrimidine (PAP) - derivatives: a new class of potent and highly selective PDGF-receptor autophosphorylation inhibitors" BIOORGANIC & MEDICINAL CHEMISTRY LETTERS, OXFORD, GB, vol. 6, no. 11, 4 June 1996 (1996-06-04), pages 1221-1226, XP004134858 ISSN: 0960-894X table 1 <div style="text-align: center; margin-top: 20px;"> --- -/-- </div>	1, 2, 15, 16
<div style="display: flex; justify-content: space-between;"> <input checked="" type="checkbox"/> Further documents are listed in the continuation of box C. <input checked="" type="checkbox"/> Patent family members are listed in annex. </div>		
<div style="display: flex;"> <div style="flex: 1;"> <p>* Special categories of cited documents:</p> <p>*A* document defining the general state of the art which is not considered to be of particular relevance</p> <p>*E* earlier document but published on or after the international filing date</p> <p>*L* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)</p> <p>*O* document referring to an oral disclosure, use, exhibition or other means</p> <p>*P* document published prior to the international filing date but later than the priority date claimed</p> </div> <div style="flex: 1;"> <p>*T* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention</p> <p>*X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone</p> <p>*Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.</p> <p>*8* document member of the same patent family</p> </div> </div>		
Date of the actual completion of the international search <div style="text-align: center; font-size: 1.2em;">3 February 2003</div>		Date of mailing of the international search report <div style="text-align: center; font-size: 1.2em;">15. 05. 03</div>
Name and mailing address of the ISA European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 340-3016		Authorized officer <div style="text-align: center; font-size: 1.2em;">Grassi, D</div>

INTERNATIONAL SEARCH REPORT

International Application No

PCT/EP 02/05420

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	ZIMMERMANN J ET AL: "Potent and selective inhibitors of the Abl-kinase: phenylamino-pyrimidine (PAP) derivatives" BIOORGANIC & MEDICINAL CHEMISTRY LETTERS, OXFORD, GB, vol. 7, no. 2, 21 January 1997 (1997-01-21), pages 187-192, XP004135990 ISSN: 0960-894X table 1 ---	1,2,15, 16
X	ZIMMERMANN J ET AL: "PHENYLAMINO-PYRIMIDINE (PAP) DERIVATIVES: A NEW CLASS OF POTENT AND SELECTIVE INHIBITORS OF PROTEIN KINASE C (PKC)" ARCHIV DER PHARMAZIE, VCH VERLAGSGESELLSCHAFT MBH, WEINHEIM, DE, vol. 329, no. 7, July 1996 (1996-07), pages 371-376, XP000885618 ISSN: 0365-6233 table I ---	1,2,15, 16
X	WO 95 09847 A (CIBA GEIGY AG ; ZIMMERMANN JUERG (CH)) 13 April 1995 (1995-04-13) cited in the application page 8, line 2; claim 1 page 13, line 1 ---	1-3,15, 16
X	EP 0 564 409 A (CIBA GEIGY AG) 6 October 1993 (1993-10-06) cited in the application claim 1 ---	1,2,15, 16
Y	JIMI T ET AL: "HIGH LEVELS OF NERVOUS SYSTEM-SPECIFIC PROTEINS IN CEREBROSPINAL FLUID IN PATIENTS WITH EARLY STAGE CRUTZFELDT-JAKOB DISEASE" CLINICA CHIMICA ACTA, AMSTERDAM, NL, vol. 211, no. 1/2, 15 October 1992 (1992-10-15), pages 37-46, XP002071132 ISSN: 0009-8981 page 37 page 40 ---	19,21, 26,42,43
Y	JAE-KWANG ET AL.: "Increased expression of CaM kinase II alpha in the brains of scrapie-infected mice" NEUROSCIENCE LETTERS, vol. 273, 1999, pages 37-40, XP002229677 abstract figure 1 page 39 ---	19,21, 26,42,43

-/--

INTERNATIONAL SEARCH REPORT

International Application No.

PCT/EP 02/05420

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 00 64894 A (KANZAKI NAUYUKI ;MIWATASHI SEIJI (JP); OHKAWA SHIGENORI (JP); TAKE) 2 November 2000 (2000-11-02) page 12, line 1-8 -& EP 1 180 518 A 20 February 2002 (2002-02-20) page 1, line 5-8 page 12, line 1-8 ---	22-25
A	US 6 107 301 A (ALDRICH PAUL EDWARD ET AL) 22 August 2000 (2000-08-22) the whole document -----	3, 10, 17, 18

INTERNATIONAL SEARCH REPORT

International application No.
PCT/EP 02/05420

Box I Observations where certain claims were found unsearchable (Continuation of Item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☒ Claims Nos.: 1, 2, 15, 16
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
see FURTHER INFORMATION sheet PCT/ISA/210
3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of Item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☒ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
3, 10, 17, 18

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Continuation of Box I.2

Claims Nos.: 1,2,15,16

The initial phase of the search revealed a very large number of documents relevant to the issue of novelty. So many documents were retrieved that it is impossible to determine which parts of the claims may be said to define subject-matter for which protection might legitimately be sought (Article 6 PCT). For these reasons, a meaningful search over the whole breadth of the independent claims 1, 2, 15, and 16 is impossible. Consequently, the search has been restricted to the use of the compounds for treating infectious diseases or neurodegenerative diseases (cf. claim 3).

The applicant's attention is drawn to the fact that claims, or parts of claims, relating to inventions in respect of which no international search report has been established need not be the subject of an international preliminary examination (Rule 66.1(e) PCT). The applicant is advised that the EPO policy when acting as an International Preliminary Examining Authority is normally not to carry out a preliminary examination on matter which has not been searched. This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

1. Claims: 3,10,17,18

Use of compounds of formula (I) for the treatment of infectious diseases or neurodegenerative diseases.

2. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein FGF-R1: Use of compounds of formula (I) as inhibitors of FGF-R1, method for detecting prion disease by detecting activity of FGF-R1, method for preventing prion disease by applying an inhibitor of FGF-R1, method for regulating production of prions by applying an inhibitor of FGF-R1, monoclonal antibody binding to FGF-R1 etc.

3. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein Tkt: Use of compounds of formula (I) as inhibitors of Tkt, method for detecting prion disease by detecting activity of Tkt, method for preventing prion disease by applying an inhibitor of Tkt, method for regulating production of prions by applying an inhibitor of Tkt, monoclonal antibody binding to Tkt etc..

4. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein Abl: Use of compounds of formula (I) as inhibitors of Abl, method for detecting prion disease by detecting activity of Abl, method for preventing prion disease by applying an inhibitor of Abl, method for regulating production of prions by applying an inhibitor of Abl, monoclonal antibody binding to Abl etc..

5. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein clk1: Use of compounds of formula (I) as inhibitors of clk1, method for detecting prion disease by detecting activity of clk1, method for preventing prion disease by applying an inhibitor of clk1, method for regulating production of prions by applying an inhibitor of clk1, monoclonal antibody binding to clk1 etc..

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

6. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein MKK7: Use of compounds of formula (I) as inhibitors of MKK7, method for detecting prion disease by detecting activity of MKK7, method for preventing prion disease by applying an inhibitor of MKK7, method for regulating production of prions by applying an inhibitor of MKK7, monoclonal antibody binding to MKK7 etc..

7. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein LIMK-2: Use of compounds of formula (I) as inhibitors of LIMK-2, method for detecting prion disease by detecting activity of LIMK-2, method for preventing prion disease by applying an inhibitor of LIMK-2, method for regulating production of prions by applying an inhibitor of LIMK-2, monoclonal antibody binding to LIMK-2 etc..

8. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein CaM-KI: Use of compounds of formula (I) as inhibitors of CaM-KI, method for detecting prion disease by detecting activity of CaM-KI, method for preventing prion disease by applying an inhibitor of CaM-KI, method for regulating production of prions by applying an inhibitor of CaM-KI, monoclonal antibody binding to CaM-KI etc..

9. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein JNK2: Use of compounds of formula (I) as inhibitors of JNK2, method for detecting prion disease by detecting activity of JNK2, method for preventing prion disease by applying an inhibitor of JNK2, method for regulating production of prions by applying an inhibitor of JNK2, monoclonal antibody binding to JNK2 etc..

10. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein CDC2: Use of compounds of

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

formula (I) as inhibitors of CDC2, method for detecting prion disease by detecting activity of CDC2, method for preventing prion disease by applying an inhibitor of CDC2, method for regulating production of prions by applying an inhibitor of CDC2, monoclonal antibody binding to CDC2 etc..

11. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein PRK: Use of compounds of formula (I) as inhibitors of PRK, method for detecting prion disease by detecting activity of PRK, method for preventing prion disease by applying an inhibitor of PRK, method for regulating production of prions by applying an inhibitor of PRK, monoclonal antibody binding to PRK etc..

12. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein PTP-SL: Use of compounds of formula (I) as inhibitors of PTP-SL, method for detecting prion disease by detecting activity of PTP-SL, method for preventing prion disease by applying an inhibitor of PTP-SL, method for regulating production of prions by applying an inhibitor of PTP-SL, monoclonal antibody binding to PTP-SL etc..

13. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein PTP-zeta: Use of compounds of formula (I) as inhibitors of PTP-zeta, method for detecting prion disease by detecting activity of PTP-zeta, method for preventing prion disease by applying an inhibitor of PTP-zeta, method for regulating production of prions by applying an inhibitor of PTP-zeta, monoclonal antibody binding to PTP-zeta etc..

14. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein HSP86: Use of compounds of formula (I) as inhibitors of HSP86, method for detecting prion disease by detecting activity of HSP86, method for preventing prion disease by applying an inhibitor of HSP86, method for regulating production of prions by applying an inhibitor of HSP86, monoclonal antibody binding to HSP86

FURTHER INFORMATION CONTINUED FROM PCT/SA/ 210

etc..

15. Claims: 13(part),19(part),21-26(part),31-37(part),
42-47(part)

Inventions related to protein GPIR-1: Use of compounds of formula (I) as inhibitors of GPIR-1, method for detecting prion disease by detecting activity of GPIR-1, method for preventing prion disease by applying an inhibitor of GPIR-1, method for regulating production of prions by applying an inhibitor of GPIR-1, monoclonal antibody binding to GPIR-1 etc..

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/EP 02/05420

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO 9509847 A	13-04-1995	AU 693475 B	02-07-1998
		AU 7697694 A	01-05-1995
		CA 2148931 A	13-04-1995
		EP 0672035 A	20-09-1995
		JP 8503971 T	30-04-1996
		US 5612340 A	18-03-1997
EP 0564409 A	06-10-1993	AT 188964 T	15-02-2000
		AU 3569493 A	07-10-1993
		BR 1100739 A	06-06-2000
		CA 2093203 A,C	04-10-1993
		CN 1077713 A,B	27-10-1993
		CZ 9300560 A	16-02-1994
		DE 59309931 D	24-02-2000
		DK 564409 T	19-06-2000
		ES 2142857 T	01-05-2000
		FI 931458 A	04-10-1993
		GR 3032927 T	31-07-2000
		HU 64050 A	29-11-1993
		IL 105264 A	11-04-1999
		JP 2706682 B	28-01-1998
		JP 6087834 A	29-03-1994
		KR 261366 B	01-08-2000
		MX 9301929 A	29-07-1994
		NO 931283 A	04-10-1993
		NZ 247299 A	26-07-1995
		PT 564409 T	30-06-2000
		RU 2125992 C	10-02-1999
		SG 43859 A	14-11-1997
		SK 28093 A	06-04-1994
		US 5521184 A	28-05-1996
		ZA 9302397 A	04-10-1993
WO 0064894 A	02-11-2000	AU 3840100 A	10-11-2000
		BR 0009952 A	26-03-2002
		CA 2370264 A	02-11-2000
		CN 1353710 T	12-06-2002
		CZ 20013805 A	17-04-2002
		EP 1180518 A	20-02-2002
		JP 3333774 B	15-10-2002
		JP 2001114779 A	24-04-2001
		JP 2002363179 A	18-12-2002
		NO 20015156 A	18-12-2001
		SK 14952001 A	04-04-2002
US 6107301 A	22-08-2000	AU 692484 B	11-06-1998
		AU 8012294 A	04-05-1995
		BR 9407799 A	06-05-1997
		CA 2174080 A	20-04-1995
		CZ 9601014 A	13-11-1996
		EP 0723533 A	31-07-1996
		HR 940664 A	31-12-1996
		HU 74464 A	30-12-1996
		NO 961425 A	12-06-1996
		NZ 274978 A	27-04-1998
		PL 313973 A	05-08-1996
		SK 47096 A	01-10-1996
		US 6342503 B	29-01-2002

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/EP 02/05420

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
US 6107301 A		CN 1142817 A	12-02-1997
		FI 961599 A	07-06-1996
		JP 9504520 T	06-05-1997
		RU 2153494 C	27-07-2000
		WO 9510506 A	20-04-1995
		ZA 9407921 A	11-04-1996
<hr/>			